

FIG.1A

2/65

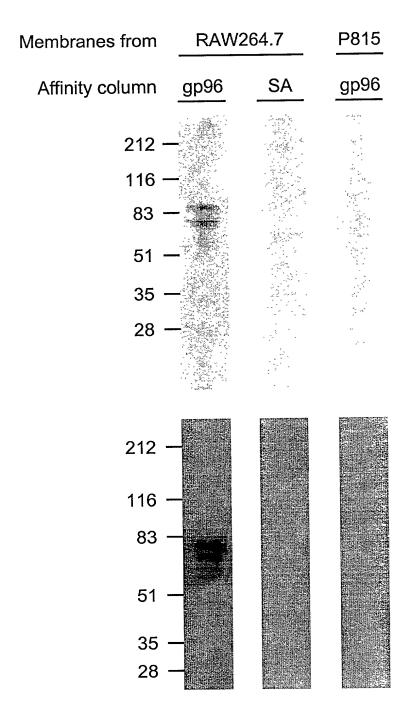


FIG.1B

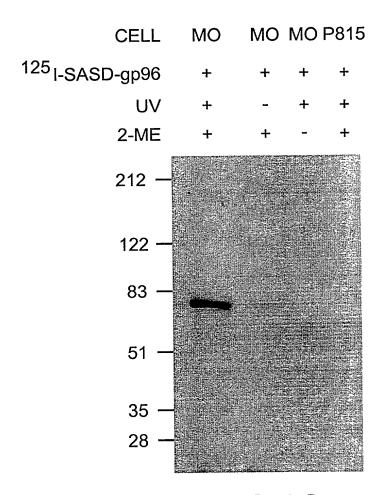


FIG.1C

	Pi	re-immur	ne	Po	st-immur	ne
	RAW264.7	Macrophage	p815	RAW264.7	Macrophage	p815
122 —						
83 —						7.9 2.4 2.5
51 —	-					eg * Nos
35 —	-	×.				

FIG.2A

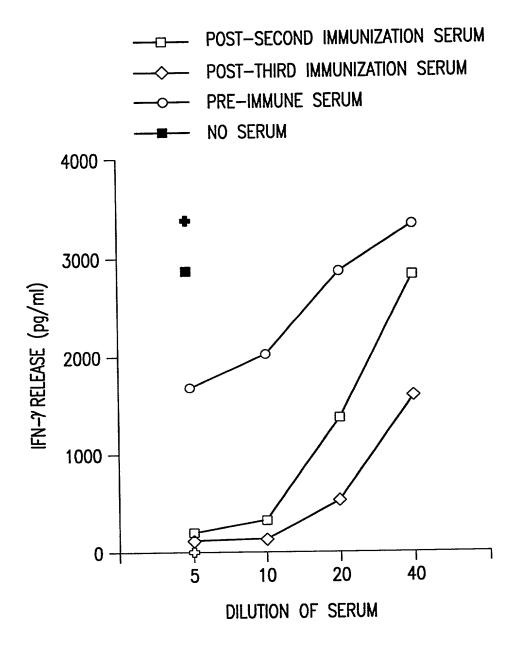


FIG.2B

Seq	#_	b	<u>y</u> _	+1
G	1	58.1	_	10
Ğ	2	115.1	1095.2	9
Ā	3	186.2	1038.2	8
Ĺ	4	299.3	967.1	7
H	5	436.5	853.9	6
1	6	549.6	716.8	5
Y	7	712.8	603.6	4
H	8	850.0	440.5	3
Q	9	978.1	303.3	2
R	10	_	175.2	1

FIG.3A

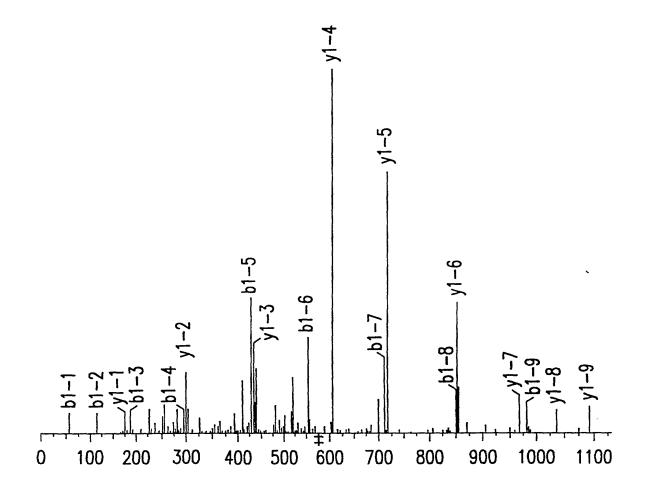


FIG.3B

POSITION	MH+	SEQUENCE				
509-518	955.0122	SGFSLGSDGK	(SEQ	ID	NO:	54)
328-337	973.1753	GIALDPAMGK	(SEQ	ID	NO:	55)
460-469	1152.3010	GGALHIYHQR	(SEQ	ID	NO:	56)
338-348	1315.5116	VFFTDYGQIPK	(SEQ	ID	NO:	57)

FIG.3C

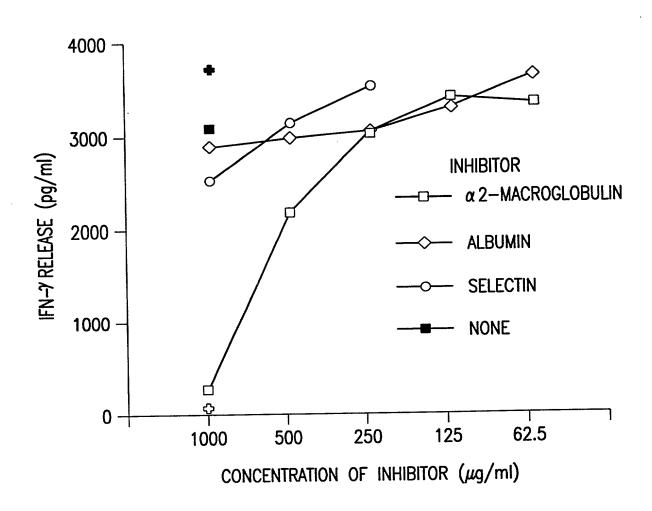
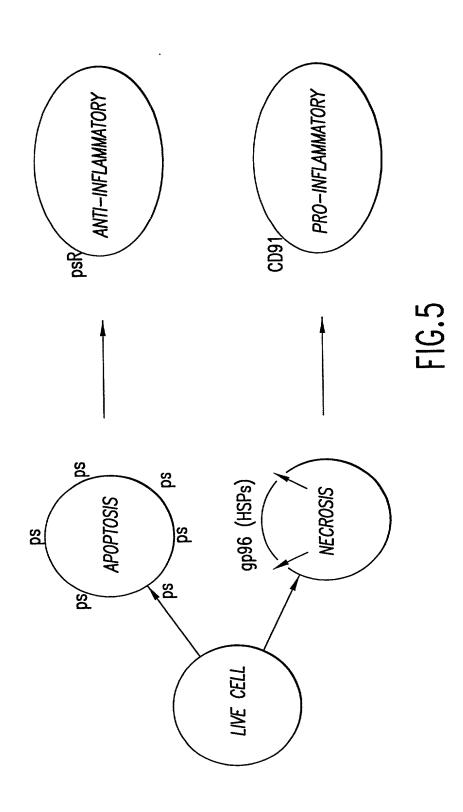


FIG.4



CGCTGCTCCC CO GGCCCCTACC A CAATTGTGCA T GAGGGGGAGA G CGCACCCGCG T CCTGGTTCGC T GGGACCCCCC A CTGTTTGCAT C	AGGCACCCC (TTTTGCAGC (GAGCGAGGA (CAGCAGGCC (TTGCTTAAG (ATTGGGGGG (ATCGGGTCC GGAGTCGGC TAAAGCAGG TTCCCAGGG AAGGATAAG GCGAGGACA ATG CTG	ACGCCCCC TCCGAGAT GGTGAAGG GGCTCGGA ATAGAAGA AGAAGTAA	CCA CCCCC TGG GGCTC GGT TCGAA AAC TGTAC AGT CGGGG ACA GGACC CCG TTG	CCACCC CO ETGAGC T ATTTGG GO CCATTT CA EAGAGG A CAGAGG G CTG CTG	GCCTCCTC FCGCCCTC GGCAGGGC ACCTATGC AGATAAAC TGGGGGGC CTC GTG	GC 120 GG 180 GG 240 CC 300 GG 360
CCG CTG CTT Pro Leu Leu	TCA GCT CTO Ser Ala Leo 15	GTC TCC Val Ser	GGG GCC A Gly Ala 1 20	ACT ATG Thr Met	Asp Ala	CCT AAA Pro Lys 25	519
ACT TGC AGC Thr Cys Ser	CCT AAG CA Pro Lys Gl 30	TTT GCC n Phe Ala	TGC AGA (Cys Arg / 35	GAC CAA Asp Gln	ATC ACC Ile Thr 40	TGT ATC Cys Ile	567
TCA AAG GGC Ser Lys Gly 45	TGG CGG TG Trp Arg Cy	T GAC GGT s Asp Gly 50	GAA AGA (Glu Arg	GAT TGC Asp Cys	CCC GAC Pro Asp 55	GGC TCT Gly Ser	615
GAT GAA GCC Asp Glu Ala 60	CCT GAG AT Pro Glu Il	C TGT CCA e Cys Pro 65	CAG AGT	AAA GCC Lys Ala 70	CAG AGA Gln Arg	TGC CCG Cys Pro	663
CCA AAT GAG Pro Asn Glu 75	CAC AGT TO His Ser Cy 80	s Leu Gly	Thr Glu	CTA TGT Leu Cys 85	GTC CCC Val Pro	ATG TCT Met Ser 90	711
CGT CTC TGC Arg Leu Cys	AAC GGG AT Asn Gly IT 95	C CAG GAC e Gln Asp	TGC ATG Cys Met 100	GAT GGC Asp Gly	TCA GAC Ser Asp	GAG GG Glu Gly 105	Г 759 У
GCT CAC TGC Ala His Cys	CGA GAG C Arg Glu Lo	C CGA GCC u Arg Ala	C AAC TGT A Asn Cys 115	TCT CGA Ser Arg	ATG GGT Met Gly 120	Cys Gl	A 807 n
CAC CAT TGT His His Cys 125	: Val Pro T	CA CCC AGT or Pro Ser 130	r Gly Pro	ACG TGC Thr Cys	TAC TGT Tyr Cys 135	AAC AG Asn Se	C 855 r

12/65 AGC TTC CAG CTC GAG GCA GAT GGC AAG ACG TGC AAA GAT TTT GAC GAG Ser Phe Gln Leu Glu Ala Asp Gly Lys Thr Cys Lys Asp Phe Asp Glu TGT TCC GTG TAT GGC ACC TGC AGC CAG CTT TGC ACC AAC ACA GAT GGC Cys Ser Val Tyr Gly Thr Cys Ser Gln Leu Cys Thr Asn Thr Asp Gly TCC TTC ACA TGT GGC TGT GTT GAA GGC TAC CTG CTG CAA CCG GAC AAC Ser Phe Thr Cys Gly Cys Val Glu Gly Tyr Leu Leu Gln Pro Asp Asn CGC TCC TGC AAG GCC AAG AAT GAG CCA GTA GAT CGG CCG CCA GTG CTA Arg Ser Cys Lys Ala Lys Asn Glu Pro Val Asp Arg Pro Pro Val Leu CTG ATT GCC AAC TCT CAG AAC ATC CTA GCT ACG TAC CTG AGT GGG GCC Leu Ile Ala Asn Ser Gln Asn Ile Leu Ala Thr Tyr Leu Ser Gly Ala CAA GTG TCT ACC ATC ACA CCC ACC AGC ACC CGA CAA ACC ACG GCC ATG Gln Val Ser Thr Ile Thr Pro Thr Ser Thr Arg Gln Thr Thr Ala Met GAC TTC AGT TAT GCC AAT GAG ACC GTA TGC TGG GTG CAC GTT GGG GAC Asp Phe Ser Tyr Ala Asn Glu Thr Val Cys Trp Val His Val Gly Asp AGT GCT GCC CAG ACA CAG CTC AAG TGT GCC CGG ATG CCT GGC CTG AAG Ser Ala Ala Gln Thr Gln Leu Lys Cys Ala Arg Met Pro Gly Leu Lys GGC TTT GTG GAT GAG CAT ACC ATC AAC ATC TCC CTC AGC CTG CAC CAC Gly Phe Val Asp Glu His Thr Ile Asn Ile Ser Leu Ser Leu His His GTG GAG CAG ATG GCA ATC GAC TGG CTG ACG GGA AAC TTC TAC TTT GTC Val Glu Gln Met Ala Ile Asp Trp Leu Thr Gly Asn Phe Tyr Phe Val GAC GAC ATT GAC GAC AGG ATC TTT GTC TGT AAC CGA AAC GGG GAC ACC Asp Asp Ile Asp Asp Arg Ile Phe Val Cys Asn Arg Asn Gly Asp Thr

FIG.6A-2

 							ATC Ile		1431
							CAG G1n 345		1479
							AAG Lys		1527
							CTG Leu		1575
							GAG G1u		1623
							ATC Ile		1671
							TAC Tyr 425		1719
 							ATC Ile	CGA Arg	1767
 	 	 			 	 -		GAC Asp	1815
 	 						CGA Arg	GTG Val	1863
					Lys			TGC Cys 490	1911

14/65 TCC GAC ATC TGC CTC CTG GCC AAC AGT CAC AAG GCA AGG ACC TGC AGG Ser Asp Ile Cys Leu Leu Ala Asn Ser His Lys Ala Arg Thr Cys Arg TGC AGG TCT GGC TTC AGC CTG GGA AGT GAT GGG AAG TCT TGT AAG AAA Cys Arg Ser Gly Phe Ser Leu Gly Ser Asp Gly Lys Ser Cys Lys Lys CCT GAA CAT GAG CTG TTC CTC GTG TAT GGC AAG GGC CGA CCA GGC ATC Pro Glu His Glu Leu Phe Leu Val Tyr Gly Lys Gly Arg Pro Gly Ile ATT AGA GGC ATG GAC ATG GGG GCC AAG GTC CCA GAT GAG CAC ATG ATC Ile Arg Gly Met Asp Met Gly Ala Lys Val Pro Asp Glu His Met Ile CCC ATC GAG AAC CTT ATG AAT CCA CGC GCT CTG GAC TTC CAC GCC GAG Pro Ile Glu Asn Leu Met Asn Pro Arg Ala Leu Asp Phe His Ala Glu ACC GGC TTC ATC TAC TTT GCT GAC ACC ACC AGC TAC CTC ATT GGC CGC Thr Gly Phe Ile Tyr Phe Ala Asp Thr Thr Ser Tyr Leu Ile Gly Arg CAG AAA ATT GAT GGC ACG GAG AGA GAG ACT ATC CTG AAG GAT GGC ATC Gln Lys Ile Asp Gly Thr Glu Arg Glu Thr Ile Leu Lys Asp Gly Ile CAC AAT GTG GAG GGC GTA GCC GTG GAC TGG ATG GGA GAC AAT CTT TAC His Asn Val Glu Gly Val Ala Val Asp Trp Met Gly Asp Asn Leu Tyr TGG ACT GAT GAT GGC CCC AAG AAG ACC ATT AGT GTG GCC AGG CTG GAG Trp Thr Asp Asp Gly Pro Lys Lys Thr Ile Ser Val Ala Arg Leu Glu AAA GCC GCT CAG ACC CGG AAG ACT CTA ATT GAG GGC AAG ATG ACA CAC Lys Ala Ala Gln Thr Arg Lys Thr Leu Ile Glu Gly Lys Met Thr His CCC AGG GCC ATT GTA GTG GAT CCA CTC AAT GGG TGG ATG TAC TGG ACA Pro Arg Ala Ile Val Val Asp Pro Leu Asn Gly Trp Met Tyr Trp Thr

FIG.6A-4

TGG Trp								2487
TGG Trp								2535
CTT Leu 700								2583
TGG Trp								2631
ACA Thr								2679
GGC G1y								2727
GGC Gly								2775
GTG Val 780							CGA Arg	2823
TAC Tyr							GTA Val 810	2871
AAC Asn							 	2919
TGT Cys								2967

	TG GCG eu Ala 845			CCC					3015
G1n Pl	TT GCC he Ala 60								3063
	GA GAC ly Asp								3111
	AT CAA is Gln								3159
	GT ATC ys Ile								3207
	GC GAG er Glu 925								3255
Pro A	AC CAG sn G1n 40								3303
	GT GAT ys Asp								3351
	GC GCC ys Ala								 3399
	GC AGA ily Arg						Asn	 	 3447
	GG GAC ly Asp 1005		Glu			His			 3495

FIG.6A-6

								17/	65							
ACC (Thr (CAG G1n O20	TTC Phe	AAG Lys	TGC Cys	Asn	AGT (Ser (025	GGC Gly	AGA Arg	TGC / Cys	Ile	CCC Pro .030	GAG G1u	CAC His	TGG Trp	ACG Thr	3543
TGT (Cys / 1035	GAT Asp	GGG Gly	GAC Asp	Asn	GAT Asp L040	TGT Cys	GGG Gly	GAC Asp	Tyr	AGC Ser 045	GAC Asp	GAG G1u	ACA Thr	His	GCC A1 a .050	3591
AAC Asn	TGT Cys	ACC Thr	Asn	CAG G1n L055	GCT Ala	ACA Thr	AGA Arg	Pro	CCT Pro .060	GGT Gly	GGC Gly	TGC Cys	His	TCG Ser 1065	GAT Asp	3639
GAG i Glu i	TTC Phe	Gln	TGC Cys 1070	CCG Pro	CTA Leu	GAT Asp	G1y	CTG Leu L075	TGC Cys	ATC Ile	CCC Pro	Leu	AGG Arg 1080	TGG Trp	CGC Arg	3687
TGC Cys	Asp	GGG Gly 1085	GAC Asp	ACC Thr	GAC Asp	Cys	ATG Met L090	GAT Asp	TCC Ser	AGC Ser	Asp	GAG Glu 1095	Lys	AGC Ser	TGT Cys	3735
Glu	GGC Gly .100	GTG Val	ACC Thr	CAT His	Val	TGT Cys 1105	GAC Asp	CCG Pro	AAT Asn	Val	AAG Lys 1110	Phe	GGC Gly	TGC Cys	AAG Lys	3783
GAC Asp 1115	TCC Ser	GCC Ala	CGG Arg	Cys	ATC Ile 1120	AGC Ser	AAG Lys	GCG Ala	Trp	GTG Val 1125	Cys	GAT Asp	GGC Gly	GAC Asp	AGC Ser 1130	3831
GAC Asp	TGT Cys	GAA G1u	GAT Asp	AAC Asn 1135	Ser	GAC Asp	Glu	Glu	Asn	Cys	G]ı	ı Ala	ı Let	GCC Ala 1145	TGC	3879
AGG Arg	CCA Pro	CCC Pro	TCC Ser 1150	His	CCC Pro	TGC Cys	GCC Ala	AAC Asn 1155	Asn	ACC Thr	C TCT Ser	GTC Val	TG0 Cys 1160	s Leu	CCT Pro	3927
CCT Pro	GAC Asp	: AAG Lys 1165	: Leu	a TGC u Cys	GAC S Asp	Gly	AAG Lys 1170	Asp	GAC Asp	TGT Cys	r GG/ s Gl:	A GA(y Ası 117!	o Gl	C TCC y Sei	G GAT Asp	3975
Glu	GG(Gly 118(/ G1	G CT(u Lei	TGT L Cys	GAC S Asp	CAG Glr 1185	Cys	TCT S Ser	CTG Leu	AA ⁻ L Asi	T AAT n Asi 119	n Gl	T GG(y Gl)	C TG y Cy:	Γ AGT s Ser	4023

4071		TGC Cys 1					Gly					Val				
4119		CAG G1n L225	Ile					Asn					Glu		••	
4167		AAC Asn		Asp					Cys					Ala		
4215		CCT Pro			Val					Cys					Ser	
4263		ATC Ile				Lys					Ser					Gly
4311		GAC Asp					Asp					Glu				
4359	His	TTC Phe 1305	Asp					Asn					Val			
4407	ATC Ile			Glu					Trp					Gln		
4455	GTG Val	GAG G1u	Phe		Thr					Asp					Arg	
4503	TGG Trp					Gly					Leu					Val
4551	GAA G1u 1370	Ile			Leu		Ser			Trp		Ile			Ala	

19/65 GTG GCC AAG CTG GAC GGA ACC CTC CGA ACC ACT CTG CTG GCG GGT GAC Val Ala Lys Leu Asp Gly Thr Leu Arg Thr Thr Leu Leu Ala Gly Asp ATT GAG CAC CCG AGG GCC ATC GCT CTG GAC CCT CGG GAT GGG ATT CTG Ile Glu His Pro Arg Ala Ile Ala Leu Asp Pro Arg Asp Gly Ile Leu TTT TGG ACA GAC TGG GAT GCC AGC CTG CCA CGA ATC GAG GCT GCA TCC Phe Trp Thr Asp Trp Asp Ala Ser Leu Pro Arg Ile Glu Ala Ala Ser ATG AGT GGA GCT GGC CGA ACC ATC CAC CGG GAG ACA GGC TCT GGG Met Ser Gly Ala Gly Arg Arg Thr Ile His Arg Glu Thr Gly Ser Gly GGC TGC GCC AAT GGG CTC ACC GTG GAT TAC CTG GAG AAG CGC ATC CTC Gly Cys Ala Asn Gly Leu Thr Val Asp Tyr Leu Glu Lys Arg Ile Leu TGG ATT GAT GCT AGG TCA GAT GCC ATC TAT TCA GCC CGG TAT GAC GGC Trp Ile Asp Ala Arg Ser Asp Ala Ile Tyr Ser Ala Arg Tyr Asp Gly TCC GGC CAC ATG GAG GTG CTT CGG GGA CAC GAG TTC CTG TCA CAC CCA Ser Gly His Met Glu Val Leu Arg Gly His Glu Phe Leu Ser His Pro TTT GCC GTG ACA CTG TAC GGT GGG GAG GTG TAC TGG ACC GAC TGG CGA Phe Ala Val Thr Leu Tyr Gly Gly Glu Val Tyr Trp Thr Asp Trp Arg ACA AAT ACA CTG GCT AAG GCC AAC AAG TGG ACT GGC CAC AAC GTC ACC Thr Asn Thr Leu Ala Lys Ala Asn Lys Trp Thr Gly His Asn Val Thr GTG GTA CAG AGG ACC AAC ACC CAG CCC TTC GAC CTG CAG GTG TAT CAC Val Val Gln Arg Thr Asn Thr Gln Pro Phe Asp Leu Gln Val Tyr His CCT TCC CGG CAG CCC ATG GCT CCA AAC CCA TGT GAG GCC AAT GGC GGC Pro Ser Arg Gln Pro Met Ala Pro Asn Pro Cys Glu Ala Asn Gly Gly

20/65 CGG GGC CCC TGT TCC CAT CTG TGC CTC ATC AAC TAC AAC CGG ACC GTC Arg Gly Pro Cys Ser His Leu Cys Leu Ile Asn Tyr Asn Arg Thr Val 1550 1555 1560 TCC TGG GCC TGT CCC CAC CTC ATG AAG CTG CAC AAG GAC AAC ACC ACC 5175

TCC TGG GCC TGT CCC CAC CTC ATG AAG CTG CAC AAG GAC AAC ACC ACC Ser Trp Ala Cys Pro His Leu Met Lys Leu His Lys Asp Asn Thr Thr

1565 1570 1575

TGC TAT GAG TTT AAG AAG TTC CTG CTG TAC GCA CGT CAG ATG GAG ATC

Cys Tyr Glu Phe Lys Lys Phe Leu Leu Tyr Ala Arg Gln Met Glu Ile

1580 1585 1590

CGG GGC GTG GAC CTG GAT GCC CCG TAC TAC AAT TAT ATC ATC TCC TTC

Arg Gly Val Asp Leu Asp Ala Pro Tyr Tyr Asn Tyr Ile Ile Ser Phe

1595 1600 1605 1610

ACG GTG CCT GAT ATC GAC AAT GTC ACG GTG CTG GAC TAT GAT GCC CGA

Thr Val Pro Asp Ile Asp Asn Val Thr Val Leu Asp Tyr Asp Ala Arg

1615

1620

1625

GAG CAG CGA GTT TAC TGG TCT GAT GTG CGG ACT CAA GCC ATC AAA AGG 5367 Glu Gln Arg Val Tyr Trp Ser Asp Val Arg Thr Gln Ala Ile Lys Arg 1630 1635 1640

GCA TTT ATC AAC GGC ACT GGC GTG GAG ACC GTT GTC TCT GCA GAC TTG 5415
Ala Phe Ile Asn Gly Thr Gly Val Glu Thr Val Val Ser Ala Asp Leu
1645 1650 1655

CCC AAC GCC CAC GGG CTG GCT GTG GAC TGG GTC TCC CGA AAT CTG TTT 5463
Pro Asn Ala His Gly Leu Ala Val Asp Trp Val Ser Arg Asn Leu Phe
1660 1665 1670

TGG ACA AGT TAC GAC ACC AAC AAG AAG CAG ATT AAC GTG GCC CGG CTG

Trp Thr Ser Tyr Asp Thr Asn Lys Lys Gln Ile Asn Val Ala Arg Leu

1675

1680

1690

GAC GGC TCC TTC AAG AAT GCG GTG GTG CAG GGC CTG GAG CAG CCC CAC

Asp Gly Ser Phe Lys Asn Ala Val Val Gln Gly Leu Glu Gln Pro His

1695

1700

1705

GGC CTG GTC GTC CAC CCG CTT CGT GGC AAG CTC TAC TGG ACT GAT GGG 5607 Gly Leu Val Val His Pro Leu Arg Gly Lys Leu Tyr Trp Thr Asp Gly 1710 1715 1720

21/65 GAC AAC ATC AGC ATG GCC AAC ATG GAT GGG AGC AAC CAC ACT CTG CTC Asp Asn Ile Ser Met Ala Asn Met Asp Gly Ser Asn His Thr Leu Leu TTC AGT GGC CAG AAG GGC CCT GTG GGG TTG GCC ATT GAC TTC CCT GAG Phe Ser Gly Gln Lys Gly Pro Val Gly Leu Ala Ile Asp Phe Pro Glu AGC AAA CTC TAC TGG ATC AGC TCT GGG AAC CAC ACA ATC AAC CGT TGC Ser Lys Leu Tyr Trp Ile Ser Ser Gly Asn His Thr Ile Asn Arg Cys AAT CTG GAT GGG AGC GAG CTG GAG GTC ATC GAC ACC ATG CGG AGC CAG Asn Leu Asp Gly Ser Glu Leu Glu Val Ile Asp Thr Met Arg Ser Gln CTG GGC AAG GCC ACT GCC CTG GCC ATC ATG GGG GAC AAG CTG TGG TGG Leu Gly Lys Ala Thr Ala Leu Ala Ile Met Gly Asp Lys Leu Trp Trp GCA GAT CAG GTG TCA GAG AAG ATG GGC ACG TGC AAC AAA GCC GAT GGC Ala Asp Gln Val Ser Glu Lys Met Gly Thr Cys Asn Lys Ala Asp Gly TCT GGG TCC GTG GTG CTG CGG AAC AGT ACC ACG TTG GTT ATG CAC ATG Ser Gly Ser Val Val Leu Arg Asn Ser Thr Thr Leu Val Met His Met AAG GTG TAT GAC GAG AGC ATC CAG CTA GAG CAT GAG GGC ACC AAC CCC Lys Val Tyr Asp Glu Ser Ile Gln Leu Glu His Glu Gly Thr Asn Pro TGC AGT GTC AAC AAC GGA GAC TGT TCC CAG CTC TGC CTG CCA ACA TCA Cys Ser Val Asn Asn Gly Asp Cys Ser Gln Leu Cys Leu Pro Thr Ser GAG ACG ACT CGC TCC TGT ATG TGT ACA GCC GGT TAC AGC CTC CGG AGC Glu Thr Thr Arg Ser Cys Met Cys Thr Ala Gly Tyr Ser Leu Arg Ser GGA CAG CAG GCC TGT GAG GGT GTG GGC TCT TTT CTC CTG TAC TCT GTA Gly Gln Gln Ala Cys Glu Gly Val Gly Ser Phe Leu Leu Tyr Ser Val

His					Gly					Pro	AAT Asn 1910					6183
				Val					Leu		GTC Val			Asp		6231
			Asn					Trp			ATG Met		Leu			6279
		Arg					Gln				GAG G1u	Asp				6327
	Gly					Glu					GAC Asp					6375
Asn					Asp					Val	ATC Ile 1990					6423
				Phe					Ile		CAG Gln			Asp		6471
			Ile					Glu			TAC Tyr		Phe			6519
		Gly					Ile					Leu			ACA Thr	6567
	Arg					Asn					•		Asn		ATC Ile	6615
Ser					Gly					Trp					ATG Met	6663

6711		GAG G1u 2					Thr					Arg				
6759	GAG G1u		Val					Met					Ser			
6807	: AAG : Lys			Gly					Arg					Ile		
6855	GGC Gly				Pro					Ala					Gly	
6903	G CAG					Phe					Lys					Ile
6951	CTC Leu 2170	Gln					Asn					Val			Gly	
6999	GGG Gly		Ala					Arg			Gly		Arg			
7047	C CTG Leu	Tyr		Ala					Ser			Asp		Ala		
7095	Γ GAG o Glu		Ser		His			Lys		Ile			Glu		Tyr	
7143	C ATG s Met			Pro		Glu			Gln		Pro			Leu		
7191	C CCG r Pro 2250	Ser			Ala		Tyr			Ala		Ala			Asn	

24/65 GGG ACC CCT AAC CGC ATC TTC TTC AGT GAC ATC CAC TTT GGG AAC ATC Gly Thr Pro Asn Arg Ile Phe Phe Ser Asp Ile His Phe Gly Asn Ile CAG CAG ATC AAT GAC GAT GGC TCG GGC AGG ACC ACC ATC GTG GAA AAT Gln Gln Ile Asn Asp Asp Gly Ser Gly Arg Thr Thr Ile Val Glu Asn GTG GGC TCT GTG GAA GGC CTG GCC TAT CAC CGT GGC TGG GAC ACA CTG Val Gly Ser Val Glu Gly Leu Ala Tyr His Arg Gly Trp Asp Thr Leu TAC TGG ACA AGC TAC ACC ACA TCC ACC ATC ACC CGC CAC ACC GTG GAC Tyr Trp Thr Ser Tyr Thr Thr Ser Thr Ile Thr Arg His Thr Val Asp CAG ACT CGC CCA GGG GCC TTC GAG AGG GAG ACA GTC ATC ACC ATG TCC Gln Thr Arg Pro Gly Ala Phe Glu Arg Glu Thr Val Ile Thr Met Ser GGA GAC GAC CAC CCG AGA GCC TTT GTG CTG GAT GAG TGC CAG AAC CTG Gly Asp Asp His Pro Arg Ala Phe Val Leu Asp Glu Cys Gln Asn Leu ATG TTC TGG ACC AAT TGG AAC GAG CTC CAT CCA AGC ATC ATG CGG GCA Met Phe Trp Thr Asn Trp Asn Glu Leu His Pro Ser Ile Met Arg Ala GCC CTA TCC GGA GCC AAC GTC CTG ACC CTC ATT GAG AAG GAC ATC CGC Ala Leu Ser Gly Ala Asn Val Leu Thr Leu Ile Glu Lys Asp Ile Arg ACG CCC AAT GGG TTG GCC ATC GAC CAC CGG GCG GAG AAG CTG TAC TTC Thr Pro Asn Gly Leu Ala Ile Asp His Arg Ala Glu Lys Leu Tyr Phe TCG GAT GCC ACC TTG GAC AAG ATC GAG CGC TGC GAG TAC GAC GGC TCC Ser Asp Ala Thr Leu Asp Lys Ile Glu Arg Cys Glu Tyr Asp Gly Ser

CAC CGC TAT GTG ATC CTA AAG TCG GAG CCC GTC CAC CCC TTT GGG TTG

His Arg Tyr Val Ile Leu Lys Ser Glu Pro Val His Pro Phe Gly Leu

		•	GAC TGG GTG CO Asp Trp Val Ai 244	rg Arg Ala	7767
g Ala Asn	Lys Tyr \		GAC ATG AAG CASP Met Lys Le		7815
			ATC GCC GTG GO Ile Ala Val A 2470		7863
r Cys Glu		Pro Cys Arg	ATC AAC AAT GO Ile Asn Asn G 2485		7911
			CAC GTC AAC TO His Val Asn C		7959
			ACC TGC CGG GG Thr Cys Arg A 25	la Val Asn	8007
s Arg Ala	Gln Asp (TGT GCC AAT GG Cys Ala Asn G 2535		8055
			TCC CAC TGC A Ser His Cys L 2550		8103
u Lys Pro		Cys Asn Ser	CGC CGC TGC A Arg Arg Cys L 2565		8151
			TCC AAC ATG C Ser Asn Met L		8199
			GAT GAG ATA CA Asp Glu Ile P 26	ro Cys Asn	8247

	Thr					Gly					CGG Arg 2					8295
Пe					Arg					Val	GAT Asp 2630					8343
				Asn					Asp		AGC Ser			Phe		8391
			Lys					G1n			GAG Glu		Thr			8439
		Ala					Cys				AAC Asn	Asp				8487
	Ser					Cys					CGC Arg					8535
Leu					Cys					Cys	ATC Ile 2710					8583
	Cys			Glu					Asn		GAG Glu			Thr		8631
			Phe					Gln			TGC Cys		Asn		Arg	8679
		Ser					Cys		Gly		GAT Asp	Asp		Gly	GAT Asp	8727
	Ser		Glu			His		Glu			Thr		Gly		TCC Ser	8775

									705							
Ser					G1y					Val				TGG Trp		8823
				Lys					Gly					GTC Val		8871
			Leu					Cys					Phe	ATG Met 2825		8919
		Arg					Lys					Asp		GAC Asp		8967
	Cys					Asp					Cys			CCA Pro		9015
Cys					Phe					G1y				AGC Ser		9063
				Cys					Asp		His			AGC Ser		9111
			Lys					Thr					Lys	TGC Cys 2905		9159
		Ser		Phe			Ser					۷a۱		Glu	GCG Ala	9207
	Leu		Asn			Asp		Cys			Gly		Asp		CGC Arg	9255
Gly		His			Glu		Leu					. Ser			AGT Ser	9303

28/65	
CAG GAC TGC GAG GAC CTC AAG ATA GGC TTT AAG TGC CGC TGT CGC CCG Gln Asp Cys Glu Asp Leu Lys Ile Gly Phe Lys Cys Arg Cys Arg Pro 2955 2960 2965 2970	9351
GGC TTC CGG CTA AAG GAC GAT GGC AGG ACC TGT GCC GAC CTG GAT GAG Gly Phe Arg Leu Lys Asp Asp Gly Arg Thr Cys Ala Asp Leu Asp Glu 2975 2980 2985	9399
TGC AGC ACC TTC CCC TGC AGC CAG CTC TGC ATC AAC ACC CAC GGA Cys Ser Thr Thr Phe Pro Cys Ser Gln Leu Cys Ile Asn Thr His Gly 2990 2995 3000	9447
AGT TAC AAG TGT CTG TGT GTG GAG GGC TAT GCA CCC CGT GGC GGT GAC Ser Tyr Lys Cys Leu Cys Val Glu Gly Tyr Ala Pro Arg Gly Gly Asp 3005 3010 3015	9495
CCC CAC AGC TGC AAA GCT GTG ACC GAT GAG GAG CCA TTT CTC ATC TTT Pro His Ser Cys Lys Ala Val Thr Asp Glu Glu Pro Phe Leu Ile Phe 3020 3025 3030	9543
GCC AAC CGG TAC TAC CTG CGG AAG CTC AAC CTG GAC GGC TCC AAC TAC Ala Asn Arg Tyr Tyr Leu Arg Lys Leu Asn Leu Asp Gly Ser Asn Tyr 3035 3040 3045 3050	9591
ACA CTG CTT AAG CAG GGC CTG AAC AAT GCG GTC GCC TTG GCA TTT GAC Thr Leu Leu`Lys Gln Gly Leu Asn Asn Ala Val Ala Leu Ala Phe Asp 3055 3060 3065	9639
TAC CGA GAG CAG ATG ATC TAC TGG ACG GGC GTG ACC ACC CAG GGC AGC Tyr Arg Glu Gln Met Ile Tyr Trp Thr Gly Val Thr Thr Gln Gly Ser 3070 3075 3080	9687
ATG ATT CGC AGG ATG CAC CTC AAC GGC AGC AAC GTG CAG GTT CTG CAC Met Ile Arg Arg Met His Leu Asn Gly Ser Asn Val Gln Val Leu His 3085 3090 3095	9735
CGG ACG GGC CTT AGT AAC CCA GAT GGG CTC GCT GTG GAC TGG GTG GGT Arg Thr Gly Leu Ser Asn Pro Asp Gly Leu Ala Val Asp Trp Val Gly 3100 3105 3110	9783
GGC AAC CTG TAC TGG TGT GAC AAG GGC AGA GAT ACC ATT GAG GTG TCC Gly Asn Leu Tyr Trp Cys Asp Lys Gly Arg Asp Thr Ile Glu Val Ser 3115 3120 3125 3130	9831

29/65 AAG CTT AAC GGG GCC TAT CGG ACA GTG CTG GTC AGC TCT GGC CTC CGG Lys Leu Asn Gly Ala Tyr Arg Thr Val Leu Val Ser Ser Gly Leu Arg GAG CCC AGA GCT CTG GTA GTG GAT GTA CAG AAT GGG TAC CTG TAC TGG Glu Pro Arg Ala Leu Val Val Asp Val Gln Asn Gly Tyr Leu Tyr Trp ACA GAC TGG GGT GAC CAC TCA CTG ATC GGC CGG ATT GGC ATG GAT GGA Thr Asp Trp Gly Asp His Ser Leu Ile Gly Arg Ile Gly Met Asp Gly TCT GGC CGC AGC ATC ATC GTG GAC ACT AAG ATC ACA TGG CCC AAT GGC Ser Gly Arg Ser Ile Ile Val Asp Thr Lys Ile Thr Trp Pro Asn Gly CTG ACC GTG GAC TAC GTC ACG GAA CGC ATC TAC TGG GCT GAC GCC CGT Leu Thr Val Asp Tyr Val Thr Glu Arg Ile Tyr Trp Ala Asp Ala Arg GAG GAC TAC ATC GAG TTC GCC AGC CTG GAT GGC TCC AAC CGT CAC GTT Glu Asp Tyr Ile Glu Phe Ala Ser Leu Asp Gly Ser Asn Arg His Val GTG CTG AGC CAA GAC ATC CCA CAC ATC TTT GCG CTG ACC CTA TTT GAA Val Leu Ser Gln Asp Ile Pro His Ile Phe Ala Leu Thr Leu Phe Glu GAC TAC GTC TAC TGG ACA GAC TGG GAA ACG AAG TCC ATC AAC CGG GCC Asp Tyr Val Tyr Trp Thr Asp Trp Glu Thr Lys Ser Ile Asn Arg Ala CAC AAG ACC ACG GGT GCC AAC AAA ACA CTC CTC ATC AGC ACC CTG CAC His Lys Thr Thr Gly Ala Asn Lys Thr Leu Leu Ile Ser Thr Leu His CGG CCC ATG GAC TTA CAT GTA TTC CAC GCC CTG CGC CAG CCA GAT GTG Arg Pro Met Asp Leu His Val Phe His Ala Leu Arg Gln Pro Asp Val

FIG.6A-19

CCC AAT CAC CCC TGC AAA GTC AAC AAT GGT GGC TGC AGC AAC CTG TGC

Pro Asn His Pro Cys Lys Val Asn Asn Gly Gly Cys Ser Asn Leu Cys

								30	/b5						
		Ser					His					Pro		AAC Asn	10407
	Leu					Arg					Asn			GCA Ala	10455
Gln					Asn					Pro				AAG Lys	10503
				Asp					Ser					GAC Asp	10551
			Lys					Gln					Thr	GGC G1y 3385	10599
		Asn					Cys					Asp		CAA Gln	10647
	Ser					Cys					Cys			AGC Ser	10695
Phe					Thr					Pro				CGT Arg	10743
				Asn					Glu					TGC Cys	10791
			Cys					Phe					Thr	AAG Lys 3465	 10839
		Pro					Cys					His		GTG Val	10887

					000		TOT	100	,00	• ТО		TOT	004	OTO	OAT	10005
	Ser	GAT Asp 3485				Asn					Thr					10935
Glu		CGC Arg			Asp					Ile						10983
		GGA Gly		Asp					Gly					Lys		11031
		GAT Asp	G1u					Pro					Cys			11079
		TGT Cys					Trp					Asp				11127
	Asp	AAC Asn 3565				Glu					Arg					11175
Ser		TTT Phe			Ala					Ile						11223
		GGG Gly		His					Gly					Asp		11271
		CGC Arg	Cys					Phe					Gly			11319
		Leu					Asp					Cys		Asp	GGC Gly	11367
	Asp					Gly					Thr		Pro		GAT Asp	11415

									/00							
Glu					AAC Asn					Pro						11463
				Asp	TGT Cys 3680				Ser					Glu		11511
			Phe		TGC Cys			Asn					Cys			11559
		Val			TGG Trp		Gly					Gly				11607
	Gly				GAC Asp	Glu					Pro					11655
Asn					GAC Asp					Leu						11703
				Ser	CTG Leu 3760				Met					Gly		11751
			Glu		GAT Asp			Ile					Thr			11799
		Asn			ATG Met		Gly					Cys			ACT Thr	11847
	Lys				TGT Cys	Ala					Phe		Thr		CCG Pro	11895
Gly					Gln					Cys					ACC Thr	11943

								33	(00							
				Trp				AAG Lys	Gly					Ser		11991
			Phe					AAC Asn 3					Glu			12039
		Gln					Ala	GAT Asp 8875				Ile				12087
	Pro					Ser		TAC Tyr			Thr					12135
G1u					Asp			GAT Asp		His						12183
				Asn				GGC Gly	Thr					Ser		12231
			Ala					TCC Ser					Arg			12279
		Gly					Asn	ATT Ile 3955				Lys				12327
	Ile					Val		GGG Gly			Tyr				TCC Ser	12375
Gly					Glu			CAA G1n		Lys					AAG Lys	12423
				Gly					Pro		Ala			Val	GAC Asp 4010	12471

									700							
			G1y					Ser					CAC His			12519
		Thr					Gly					Thr	CTC Leu 4040			12567
	Asn					Thr					Asp		CAC His			12615
Arg					Asp					Va1			AGC Ser			12663
				Asp					Ala				CGA Arg	Gly		12711
			Phe					Phe					TAC Tyr			12759
		Ile					Phe					Phe	GGA Gly 4120			12807
	Leu					Gly					Ala		GAT Asp			12855
Leu					Lys					Thr		Pro	TGT Cys		CGC Arg	12903
				Trp					Ser		Ser		CCT Pro	Val	TGC Cys 4170	12951
			Asn		Lys			Asp					Val			12999

		CCA						GCC								13047
Pro	Ser	Pro 4	Thr 190	Pro	Pro	Pro		Ala 195	Pro	Arg	Pro		Thr 200	Cys	Thr	
		TGC Cys														13095
		1205					210					215	J			
		TGC Cys														13143
	1220	-3-	· J	-0 -		225	J	J			1230		•			
		TGC Cys														13191
4235		- 3			1240	J				1245					4250	
		GGC Gly														13239
				1255		·			1260		-			4265		
		ACC Thr														13287
·	·		1270			-		4275					4280			
															GGC Gly	13335
J		4285			·		4290			-		4295				
															GAG Glu	13383
	4300	v	•	J		4305	·				4310			-		
															CGC Arg	13431
4315		J		-	4320				-	4325					4330	
															AGT Ser	13479
Q			_	4335		J			4340					4345		
		CTC Leu													GTC Val	13527
J	•		4350	-		•		4355		J			4360	•		

	Cys	AAC Asn				Gly	Arg	GTA			Ser	Cys				13575
Ile	GAT	CAC His			Asn	GGT				Thr	ATG					13623
ATG	CCT	GAG Glu		Gln	TGC	CCG			Met	ACA	GGA			Cys		13671
		GTT Val	Val					Pro					Ser			13719
		CTG Leu					Leu					Ala				13767
	Trp	TAT Tyr 4445				Val					Gly					13815
Arg		ACC Thr			Ala					Ile						13863
	Met	TAT Tyr		Gly		Glu			Asp		Gly			Leu	GAT Asp 4490	13911
			Ala					Lys		Thr			Thr		CCA Pro	13959
		Ala					Gly		His			Arg		Ser	CTG Leu	14007
	Ser					Arg		Leu			' Arg		Pro		GAC Asp	14055

37/65 GAG ATA GGA GAT CCC TTG GCA TAGGGCCCTG CCCCGACGGA TGTCCCCAGA AAGC CCCCTGCCAC ATGAGTCTTT CAATGAACCC CCTCCCCAGC CGGCCCTTCT CCGGCCCTGC Glu Ile Gly Asp Pro Leu Ala 4540 4545	14110 14170
CGGGTGTACA AATGTAAAAA TGAAGGAATT ACTTTTTATA TGTGAGCGAG CAAGCGAGCA	14230
AGCACAGTAT TATCTCTTTG CATTTCCTTC CTGCCTGCTC CTCAGTATCC CCCCCATGCT	14290
GCCTTGAGGG GGCGGGGAGG GCTTTGTGGC TCAAAGGTAT GAAGGAGTCC ACATGTTCCC	14350
TACCGAGCAT ACCCCTGGAA GCCTGGCGGC ACGGCCTCCC CACCACGCCT GTGCAAGACA	14410
CTCAACGGGG CTCCGTGTCC CAGCTTTCCT TTCCTTGGCT CTCTGGGGTT AGTTCAGGGG	14470
AGGTGGAGTC CTCTGCTGAC CCTGTCTGGA AGATTTGGCT CTAGCTGAGG AAGGAGTCTT	14530
TTAGTTGAGG GAAGTCACCC CAAACCCCAG CTCCCACTTT CAGGGGCACC TCTCAGATGG	14590
CCATGCTCAG TATCCCTTCC AGACAGGCCC TCCCCTCTCT AGCGCCCCCT CTGTGGCTCC	14650
TAGGGCTGAA CACATTCTTT GGTAACTGTC CCCCAAGCCT CCCATCCCCC TGAGGGCCAG	14710
GAAGAGTCGG GGCACACCAA GGAAGGGCAA GCGGGCAGCC CCATTTTGGG GACGTGAACG	14770
TITTAATAAT TITTGCTGAA TTCCTTTACA ACTAAATAAC ACAGATATTG TTATAAATAA	14830
AATTGTAAAA AAAAAAAA	

FIG.6A-27

Met 1	Leu	Thr	Pro	Pro 5	Leu	Leu	Leu	Leu	Val 10	Pro	Leu	Leu	Ser	Ala 15	Leu
			20		Met			25					30		
		35			Gln		40					45			
•	50				Cys	55					60				
65					Ala 70					75					80
				85	Cys				90					95	
			100		Gly			105					110		
		115			Arg		120					125			
	130				Cys	135					140				
145					Lys 150					155					160
				165					170					175	
			180					185					190		Lys
		195		_			200					205			Gln
	210					215					220				Thr
225					230					235)				240
				245	•				250)				255	
			260)				265)				270)	ı His
Thr	Ile	Asr 275		Ser	Leu	Ser	Leu 280		His	s Val	Glu	G1r 285		: Ala	a Ile
Ť	290)				295	· •				300)			Arg
Ile 305		e Val	Cys	Asr	310		ı Gly	/ Asp	Thr	Cys 315		Thr	· Le	ı Lei	320

FIG.6B-1

									_	_	_	_			
Leu	Glu	Leu	Tyr	Asn 325	Pro	Lys	Gly		A1a 330	Leu	Asp	Pro	Ala	Met 335	Gly
Lys	Val	Phe	Phe 340	Thr	Asp	Tyr		G1n 345	Ile	Pro	Lys	Val	G1u 350	Arg	Cys
Asp	Met	Asp 355		Gln	Asn	Arg	Thr 360	Lys	Leu	Val	Asp	Ser 365	Lys	Ile	Val
Phe	Pro 370		Gly	Ile	Thr	Leu 375		Leu	Val	Ser	Arg 380		Val	Tyr	Trp
A1a 385		Ala	Tyr	Leu	Asp 390		Ile	Glu	Val	Val 395		Tyr	Glu	Gly	Lys 400
Gly	Arg	G1n	Thr	Ile 405		Gln	Gly	Ile	Leu 410		Glu	His	Leu	Tyr 415	
Leu	Thr	Val	Phe 420		Asn	Tyr	Leu	Tyr 425		Thr	Asn	Ser	Asp 430	Asn	Ala
Asn	Thr	G1n 435		Lys	Thr	Ser	Va1 440		Arg	Val	Asn	Arg 445	Phe	Asn	Ser
Thr	G1u 450		Gln	Val	Val	Thr 455		Val	Asp	Lys	Gly 460	Gly	Ala	Leu	His
Ile 465		His	Gln	Arg	Arg 470		Pro	Arg	Val	Arg 475		His	Ala	Cys	G1u 480
	Asp	Gln	Tyr	Gly 485		Pro	Gly	Gly	Cys 490		Asp	Ile	Cys	Leu 495	
Ala	Asn	Ser	His 500	Lys		Arg	Thr	Cys 505	Arg	Cys	Arg	Ser	Gly 510		Ser
Leu	Gly	Ser 515	Asp		Lys	Ser	Cys 520		Lys	Pro	Glu	His 525		Leu	Phe
Leu	Va1 530			Lys	Gly	Arg 535		Gly	Ile	Ile	Arg 540		Met	Asp	Met
Gly 545	Ala	Lys	Val	Pro	Asp 550		His	Met	Ile	Pro 555		Glu	Asn	Leu	Met 560
		Arg	Ala	Leu 565		Phe	His	Ala	G1u 570		Gly	Phe	: Ile	Tyr 575	Phe
Ala	Asp	Thr	Thr 580		Tyr	Leu	Ile	Gly 585		Glr	Lys	Ile	Asp 590		Thr
Glu	Arg	G1u 595		Ile	Leu	Lys	Asp 600		Ile	His	Asn	Va1 605		Gly	' Val
Ala	Val 610		Trp	Met	: Gly	Asp 615		Leu	Tyr	Trp	Thr 620) Asp	Gly	Pro
Lys 625	-	Thr	· Ile	Ser	Val 630		Arg	Leu	Glu	Lys 635		ı Ala	a G1r	1 Thr	Arg 640
Lys	Thr	Leu	ı Ile	GTu 645		' Lys	Met	Thr	His 650		Arg	y Ala	a Ile	e Val 655	Val

Asp Pro Leu Asn Gly Trp Met Tyr Trp Thr Asp Trp Glu Asp Glu Asp Asp Trp Met Asp Gly Asp Gly Asp Leu Glu Arg Ala Trp Met Asp Gly Asp Gly Asp Gly Asp Gly Arg Leu Arg Leu Trp Pro Asp Arg Leu Arg Leu Trp Pro Asp Arg Leu Arg Leu Trp Pro Asp Arg Leu Arg Leu Arg Lys Arg L	Ser Gly Phe 720 Ile His
His Arg Asp Ile Phe Val Thr Ser Lys Thr Val Leu Trp Pro Asn 690 695 700 Leu Ser Leu Asp Ile Pro Ala Gly Arg Leu Tyr Trp Val Asp Ala 705 710 715 715 715 725 730 735 735 735 740 745 745 750 685	Gly Phe 720 Ile His
His Arg Asp Ile Phe Val Thr Ser Lys Thr Val Leu Trp Pro Asn 690 695 700 Leu Ser Leu Asp Ile Pro Ala Gly Arg Leu Tyr Trp Val Asp Ala 705 710 715 Tyr Asp Arg Ile Glu Thr Ile Leu Leu Asn Gly Thr Asp Arg Lys 725 730 735 Val Tyr Glu Gly Pro Glu Leu Asn His Ala Phe Gly Leu Cys His 740 745 750	Phe 720 Ile His
Leu Ser Leu Asp Ile Pro Ala Gly Arg Leu Tyr Trp Val Asp Ala 705 Tyr Asp Arg Ile Glu Thr Ile Leu Leu Asn Gly Thr Asp Arg Lys 725 Val Tyr Glu Gly Pro Glu Leu Asn His Ala Phe Gly Leu Cys His 740 740 750 760 775 775	/20 Ile His Arg
Tyr Asp Arg Ile Glu Thr Ile Leu Leu Asn Gly Thr Asp Arg Lys 725 730 735 Val Tyr Glu Gly Pro Glu Leu Asn His Ala Phe Gly Leu Cys His 740 745 750	His Arg
Val Tyr Glu Gly Pro Glu Leu Asn His Ala Phe Gly Leu Cys His 740 745 750	Arg
770	
Gly Asn Tyr Leu Phe Trp Thr Glu Tyr Arg Ser Gly Ser Val Tyr	Arg
Leu Glu Arg Gly Val Ala Gly Ala Pro Pro Thr Val Thr Leu Leu	
Ser Glu Arg Pro Pro Ile Phe Glu Ile Arg Met Tyr Asp Ala His	G1u 800
Gln Gln Val Gly Thr Asn Lys Cys Arg Val Asn Asn Gly Gly Cys	Ser
Ser Leu Cys Leu Ala Thr Pro Gly Ser Arg Gln Cys Ala Cys Ala	
Asp Gln Val Leu Asp Thr Asp Gly Val Thr Cys Leu Ala Asn Pro	Ser
Tyr Val Pro Pro Pro Gln Cys Gln Pro Gly Gln Phe Ala Cys Ala	Asn
Asn Arg Cys Ile Gln Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp	Cys 880
Leu Asp Asn Ser Asp Glu Ala Pro Ala Leu Cys His Gln His Thr	Cys
Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asr	
Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu	ı Ser
915 920 925 Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser	· Cys
930 935 940 Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp	
945 950 955 Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro	960 Thr
965 970 975 Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile 980 985 990	

FIG.6B-3

		995					000				1	.005			
1	010				1	Ser .015]	L020				
Ser 025	Gly			1	.030	Glu]	L035				1	.040
Cys	Gly	Asp		Ser L045		Glu		His	Ala .050	Asn	Cys	Thr	Asn 1	G1n 1055	Ala
Thr	Arg		Pro 1060	Gly	Gly	Cys	His	Ser 1065	Asp	Glu	Phe	Gln	Cys L070	Pro	Leu
Asp		Leu 1075	Cys	Ile	Pro	Leu 1	Arg L080	Trp	Arg	Cys	Asp	Gly 1085	Asp	Thr	Asp
	Met 1090	Asp	Ser	Ser		Glu 1095			Cys	Glu	Gly 1100	Val	Thr	His	Val
Cys 105	Asp	Pro	Asn		Lys 1110	Phe	Gly	Cys	Lys	Asp 1115	Ser		Arg	Cys	Ile 1120
Ser	Lys	Ala		Va1 1125		Asp	Gly	Asp	Ser 1130	Asp	Cys	Glu	Asp	Asn 1135	Ser
Asp	Glu		Asn 1140	Cys	Glu	Ala		Ala 1145			Pro	Pro	Ser 1150	His	Pro
		Asn 1155	Asn	Thr		Val	1160					1165			
	Lys 1170	Asp	Asp	Cys	Gly	Asp 1175	Gly	Ser	Asp	Glu	Gly 1180	Glu	Leu	Cys	Asp
Gln	Cys	Ser	Leu				G1y	Cys	Ser	His 1195	Asn	Cys	Ser	Val	Ala 1200
185	61.4		C1.		1190		San	Cvs	Dro			Met	Glu		Gly
				1205)				1210)				1215	
	•		1220)				1225	}				1230)	Leu
Lys	Cys	Ser 1235		Lys	Cys	Asp	Gln 1240	Asr	Lys	s Phe	e Ser	`Val 1245	Lys 5	Cys	Ser
Cys	Tyr 1250	· Glu		/ Trp	Val	Leu 1255		ı Pro) Asp	o Gly	y G1ι 1260	ı Thr)	Cys	s Arg	Ser
	ı Asp	Pro) Phe	e Lys		ı Phe		: Ile	Phe			n Arç	g His	s Glu	11e 1280
265 Arc	o a Arc	ı Ile	e Ası	. Leu	1270 Hisu		Gly	/ Asp	туі	127! r Sei		l Leu	u Va	l Pro	Gly
				1285	5				129	0				1295	5
Lei	ı Arç	g Ası	n Thi 1300		e Ala	a Leu	ı Asp	o Phe 130!		s Lei	u sei	r GII	า Sei 131(r A18)	a Leu
Tv	n Trr	ր Thi			a Va	ו ה	ı Ası			e Tv	r Ar	a G1			ı Leu
, ,,,	114	131!		- ,			1320				·	132	5		

•	Asn 1330	Gly	Ala	Leu		Ser 1335					Ile 1340	Gln	Tyr	Gly	Leu
Ala 345	Thr	Pro	Glu		Leu 1350			•	•			-		Ile 1	Tyr .360
Trp	Val	G1u		Asn 1365		Asp			G1u L370	Val	Ala	Lys		Asp l375	G1y
Thr	Leu		Thr 1380		Leu	Leu			Asp				Pro 1390	Arg	Ala
Ile		Leu 1395	-	Pro	-	-	Gly L400		Leu			Thr 1405	Asp	Trp	Asp
	Ser 1410	Leu	Pro	Arg		Glu 1415					Ser L420	Gly	Ala	Gly	Arg
Arg 425		Ile	His	_	G1u 1430			Ser	_	Gly L435	Cys	Ala	Asn	Gly	Leu L440
Thr	Val	Asp		Leu 1445		Lys			Leu 1450					Arg L455	Ser
Asp	Ala			Ser				-	Gly				Met L470	Glu	Val
Leu	_	G1y 1475	His	Glu	Phe		Ser 1480		Pro			Va1 1485	Thr	Leu	Tyr
				Tyr	-	Thr 1495	-	Trp			Asn 1500	Thr	Leu	Ala	Lys
A1a 505	Asn	Lys	Trp		Gly 1510			Val		Va1 1515	Val	Gln	Arg	Thr	Asn 1520
Thr	Gln	Pro		Asp 1525					His 1530		Ser			Pro 1535	Met
			1540					1545					1550	Ser	
		1555					1560					1565		Pro	
	Met 1570	Lys	Leu	His	-	Asp 1575	Asn	Thr	Thr	-	Tyr 1580	Glu	Phe	Lys	Lys
Phe 585	Leu	Leu	Tyr		Arg 1590	Gln	Met	Glu		Arg 1595	Gly	Val	Asp	Leu	Asp 1600
Ala	Pro	Tyr		Asn 1605	Tyr	Ile	Ile		Phe 1610	Thr	Val	Pro	-	Ile 1615	Asp
Asn	Va1		Va1 1620	Leu	Asp	Tyr	-	Ala 1625	Arg	Glu	Gln		Val 1630	Tyr	Trp
Ser	•	Val 1635	Arg	Thr	Gln		Ile 1640	Lys	Arg	Ala				Gly	Thr
	Val 1650	Glu	Thr	Val		Ser 1655	Ala	Asp	Leu				His	Gly	Leu

A1a 665	Val	Asp	Trp				Asn			Trp 1675			Tyr	Asp 1	Thr .680
Asn	Lys	Lys			Asn	Val		Arg						Lys .695	Asn
Ala	Va1		G1n 1700	Gly	Leu	Glu			His				Val 1710	His	Pro
Leu			Lys			-		Asp		Asp	Asn	I1e l725	Ser	Met	Ala
]	L730	·			1	1735				1	L740			Lys	
745				-	1750					L755					1760
			1	L765				1	L770			•	1	Ser 1775	
]	L780]	1785				:	1790	Thr	
]	1795			-		1800	·				1805		Ser	
-	1810				-	1815		·		:	1820			Val	
825				:	1830					1835			·		L840
			1	L845					1850					Asn 1855	
		1	1860					1865					1870	Ser	
•	-	1875				•	1880				•	1885		Cys	
	1890					1895					1900			Arg	_
905					1910					1915					1920
				1925					1930					Asp 1935	
			1940					1945					1950	Lys	
		1955					1960					1965		Arg	
	1970					1975					1980			Thr	·
GIN 985	Gly	rne	Asp		11e 1990	Glu	Vai	Ala		Leu 1995	Asn	Gly	Ser	Phe	Arg 2000

Tyr	Val	Val		Ser 2005	Gln	Gly	Leu	•	-	Pro	_	Ala		Thr 1015	Val
His	Pro		-	-	Tyr			Trp 2025		Glu	•	Gly 2	His 2030	Tyr	Pro
Arg		G1u 2035				Leu						Va1 2045	Val	Leu	Val
	Va1 2050	Ser	Ile	Ser		Pro 2055	Asn	Gly	Ile	Ser	Va1 2060	Asp	Tyr	G1n	Gly
G1 y 065	Lys	Leu	Tyr		Cys 2070					•	Lys	Ile	Glu		Ile 2080
Asp	Leu	Glu		Gly 2085	Glu	Asn	Arg	Glu				Ser		Asn 2095	Asn
Met	Asp		Phe 2100	Ser	Va1	Ser		Phe 2105	Glu	Asp	Phe	Ile 2	Tyr 2110	Trp	Ser
Asp	_	Thr 2115	His		Asn			Ile		Arg		Cys 2125	Lys	Asp	Asn
	Thr 2130	Asp	Ser	Val		Leu 2135	Arg	Thr	Gly		Gly 2140	Val	Gln	Leu	Lys
Asp 145	Ile	Lys	Val		Asn 2150	Arg	Asp	Arg		Lys 2155	Gly	Thr	Asn		Cys 2160
Ala	Val	Ala		Gly 2165		Cys	Gln		Leu 2170	Cys		Tyr		G1y 2175	Gly
Gly	Gln		Ala 2180		Ala							Ala 2	G1u 2190	Asp	G1y
Ala			Arg		Tyr			Tyr		Leu		Ser 2205	Glu	Arg	Thr
	Leu 2210	Lys	Ser	Ile		Leu 2215	Ser	Asp	Glu	-	Asn 2220	Leu	Asn	Ala	Pro
Val 225	G1n	Pro	Phe		Asp 2230	Pro		His		Lys 2235	Asn	Val	Ile		Leu 2240
Ala	Phe	Asp		Arg 2245	Ala	Gly	Thr		Pro 2250	Gly	Thr	Pro		Arg 2255	Ile
Phe	Phe		Asp 2260	Ile	His	Phe		Asn 2265	Ile	Gln	Gln		Asn 2270	Asp	Asp
Gly		G1y 2275	Arg	Thr	Thr		Val 2280	Glu	Asn	Val		Ser 2285	Val	G1u	Gly
	Ala 2290	Tyr	His	Arg		Trp 2295	Asp	Thr	Leu		Trp 2300	Thr	Ser	Tyr	Thr
Thr 305	Ser	Thr	Ile		Arg 2310	His	Thr	Val		G1n 2315	Thr	Arg	Pro		A1a 2320
Phe	Glu	Arg		Thr 2325	Val	Ile	Thr				Asp	Asp			Arg

Ala	Phe		Leu 2340	Asp	Glu	Cys		Asn 2345		Met	Phe		Thr 2350	Asn	Trp
Asn		Leu 2355	His	Pro	Ser			Arg		Ala		Ser 2365	Gly	Ala	Asn
	Leu 2370	Thr	Leu	Ile		Lys 2375	Asp	Ile	Arg		Pro 2380	Asn	Gly	Leu	Ala
Ile 385	Asp	His	Arg		G1u 2390			Tyr		Ser 2395	Asp	Ala	Thr		Asp 2400
Lys	Ile	Glu	Arg	Cys 2405	Glu	Tyr	Asp		Ser 2410	His	Arg	Tyr		Ile 2415	Leu
Lys	Ser		Pro 2420		His	Pro	Phe	Gly					Gly 2430	Glu	His
Ile		Trp 2435	Thr		Trp		Arg 2440		Ala			Arg 2445	Ala	Asn	Lys
	Va1 2450	Gly	Ser	Asp		Lys 2455		Leu	Arg		Asp 2460	Ile	Pro	Gln	Gln
Pro 465	Met	Gly	Ile		A1a 2470	Val		Asn	•		Asn		Cys		Leu 2480
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Cys 625	Asn	Gln	Phe		Asp 2630	Cys	Glu	Asp		Ser 2635	Asp	Glu	Met		Cys 2640
Ser	Ala	Thr	Asp	Cys 2645	Ser	Ser	Tyr		Arg 2650	Leu	Gly	Val		Gly 2655	Va1
Leu	Phe		Pro 2660	Cys	Glu	Arg		Ser 2665	Leu	Cys	Tyr		Pro 2670	Ser	Trp

Val		Asp 2675	Gly	Ala	Asn		Cys 2680	Gly	Asp	Tyr		Asp 2685	Glu	Arg	Asp
	Pro 2690	Gly	Val	Lys		Pro 2695	Arg		Pro	Leu 2	Asn 2700	Tyr	Phe	Ala	Cys
Pro 705	Ser	G1y	Arg		Ile 2710		Met		-	Thr 2715	Cys	Asp	Lys		Asp 2720
Asp	Cys	Glu		G1y 2725	Glu	•	Glu			Cys		Lys		Cys 2735	Ser
Glu	Ala		Phe 2740	Glu	Cys	Gln		His 2745	Arg	Cys	Ile		Lys 2750	G1n	Trp
Leu		Asp 2755		Ser	Asp		Cys 2760	Gly.	Asp	Gly		Asp 2765	Glu	Ala	Ala
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Thr 785	His	Va1	Cys		Pro 2790			Trp		Cys 2795	Asp	Gly	Asp		Asp 2800
Cys	Thr	Asp	_	A1a 2805	Asp	Glu	Ser		Thr 2810	Ala		Cys		Tyr 2815	Asn
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Pro		His 2835	Phe	Val	Cys		His 2840	Asp	Arg	Asp		A1a 2845	Asp	Gly	Ser
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Arg 865	Cys	Ala	Asn		Arg 2870	Cys	Leu	Ser		Arg 2875	Gln	Trp	Glu		Asp 2880
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Cys 945	Leu	Ser	Arg		Leu 2950	Ser	Gly	Cys		G1n 2955	Asp	Cys	Glu	-	Leu 2960
Lys	Ile	Gly		Lys 2965	Cys	Arg	Cys		Pro 2970	G1y	Phe	Arg		Lys 2975	Asp
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Gly Asp His Ser Asp Glu Pro Pro Asp Cys Pro Glu Phe Lys Cys Arg 3365 3370 3375 Pro Gly Gln Phe Gln Cys Ser Thr Gly IIe Cys Thr Asn Pro Ala Phe 3380 3385 3385 3390 Ile Cys Asp Gly Asp Asn Asp Cys Gln Asp Asn Ser Asp Glu Ala Asn 3395 Cys Asp Ile His Val Cys Leu Pro Ser Gln Phe Lys Cys Thr Asn Thr 3410 Asn Arg Cys IIe Pro Gly IIe Phe Arg Cys Asn Gly Gln Asp Asn Cys 425 3430 3445 Asn Arg Cys IIe Pro Gly IIe Phe Arg Cys Asn Gly Gln Asp Asn Cys 425 3430 3455 Asn Gln Phe Gln Cys Ser IIe Thr Lys Arg Cys IIe Pro Arg Val Trp 3460 Val Cys Asp Arg Asp Asn His Cys Val Asp Gly Ser Asp Glu Pro Ala 3475 3480 3485 Asn Cys Thr Gln Met Thr Cys Gly Val Asp Glu Phe Arg Cys Lys Asp 3490 3495 3500 Ser Gly Arg Cys IIe Pro Ala Arg Trp Lys Cys Asp Gly Glu Asp Asp 505 3510 S515 3520 Cys Glu Pro Tyr Gln Phe Arg Cys Lys Asn Asn Arg Cys Val Pro Gly 3540 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu 3555 Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala 3570 Asn Gly Arg Cys IIe Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 350 Ser Gly Arg Cys IIe Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 3500 Ser Gly Arg Cys IIe Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 3500 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu 3570 Asn Gly Arg Cys IIe Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 3500 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met 3600 3601 Asp Gln Phe Gln Cys Lys Ser Gly His Cys IIe Pro Leu Arg Trp Pro 3620 3620 3625 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 3675 3680	Asp 345	Lys	Cys	Ile		Phe 3350	Trp	Trp	Lys		Asp 3355	Thr	G1u	Asp	Asp 3	Cys 360
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Cys Gly Asp Gly Ser Asp Glu Pro Lys Glu Glu Cys Asp Glu Arg Thr 3525 3530 3535 Cys Glu Pro Tyr Gln Phe Arg Cys Lys Asn Asn Arg Cys Val Pro Gly 3540 3540 3545 3550 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu 3555 3560 3565 Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala 3570 3575 3580 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 3580 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 3580 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Thr Pro Arg Cys Asp Met 3605 3610 3615 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3640 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys			Thr	GIn	Met			Gly	Val	Asp			Arg	Cys	Lys	Asp
Cys Gly Asp Gly Ser Asp Glu Pro Lys Glu Glu Cys Asp Glu Arg Thr		Gly	Arg	Cys			Ala	Arg	Trp			Asp	Gly	Glu	•	•
Cys Glu Pro Tyr Gln Phe Arg Cys Lys Asn Asn Arg Cys Val Pro Gly 3540 3550 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu 3555 3560 3565 Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala 3570 3575 3580 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 585 3590 3595 3600 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met 3605 3615 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3625 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3640 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys		Gly	Asp		Ser	Asp		Pro		Glu	Glu		Asp		Arg	
3540 3545 3550 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu 3555 3560 3565 Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala 3570 3575 3580 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 585 3590 3595 3600 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met 3605 3610 3615 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3625 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3640 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	O	07	D		-			0					^			Δ1.
Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala 3570 3575 3580 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 585 3590 3595 3600 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met 3605 3610 3615 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3625 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3640 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	Lys	GIU		-	GIN	Pne	Arg			Asn	Asn	Arg			Pro	GIY
Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala 3570 3575 3580 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 585 3590 3595 3600 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met 3605 3615 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3625 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	Arg	-		Cys	Asp	Tyr			-		Gly	-		Ser	Asp	Glu
Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp 3595 3600 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met 3605 3610 3615 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3625 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys		Ser		Thr	Pro	_	Pro	Cys				Glu		Phe	Cys	A1 a
$\begin{array}{cccccccccccccccccccccccccccccccccccc$				•					-				0.7			
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		Gly	Arg	Cys			Gly	Arg	ırp			Asp	Gly	Asp		•
Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro 3620 3625 3630 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	Cys	Ala	Asp	-		Asp	Glu	Lys	•	-	Thr	Pro	Arg	-	•	Met
Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys 3635 3640 3645 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	Asp	Gln		Gln		Lys	Ser		His		Ile	Pro		Arg		Pro
Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn 3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	Cys		Ala		Ala	Asp	-	Met		Gly	Ser	•	Glu		Ala	Cys
3650 3655 3660 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys	Clv			Val	Ana	Thn			Lou	A on	٠٦.,			Cva	100	۸۵۵
			uiy	vai	arg				Leu	ASP			uin	cys	ASN	ASN
		Leu	Cys	Lys			Ala	Trp	Lys			Gly	G1 u	Asp	_	

G1y	Asp	Asn		Asp 3685	Glu	Asn	Pro		G1u 3690	Cys	Ala	Arg		Ile 8695	Cys
Pro	Pro		Arg 3700	Pro	Phe	Arg	-	_		Asp	Arg		Cys 3710	Leu	Trp
Ile		Arg 3715	Gln	Cys	Asp					Cys		Asp 3725	Gly	Thr	Asp
			Cys	Glu				Ala	Gln	Asn 3	Pro 3740	His	Cys	Lys	Asp
		G1u	Phe				Asn	G1n	_	Cys 3755		Ser	Ser		Leu 3760
Arg	Cys	Asn		Phe 3765	Asp	Asp	Cys		Asp 3770	Gly	Ser	Asp		G1u 3775	Asp
Cys	Ser		Asp 3780	Pro	Lys	Leu		Ser 3785	Cys	Ala	Thr		A1a 3790	Ser	Met
Cys		Asp 3795	Glu	Ala	Arg	•	Va1 3800			Glu		A1a 3805	Ala	Tyr	Cys
	Cys 3810	Arg	Ser	Gly		His 3815		Val		Gly	G1n 3820	Pro	Gly	Cys	G1n
Asp 825	Ile	Asn	Glu		Leu 3830	Arg	Phe	Gly		Cys 3835	Ser	Gln	Leu	•	Asn 3840
Lys	Pro	Lys		G1 y 3845	His	Leu	Cys			Ala		Asn		Met 3855	Lys
Thr	His		Thr 3860	Cys	Lys	Ala		G1y 3865	Ser	Glu	Tyr		Va1 3870	Leu	Tyr
Пe		Asp 3875	Asp	Asn	Glu		Arg 3880			Phe		G1 y 3885	His	Pro	His
	A1a 3890	Tyr	Glu	Gln		Phe 3895		Gly		Glu 3	Ser 3900	Val	Arg	Ile	Asp
Ala 905	Met	Asp	Val		Val 3910	Lys	Ala	Gly		Val 3915	Tyr	Trp	Thr		Trp 3920
His	Thr	Gly		Ile 3925	Ser	Tyr	Arg		Leu 3930	Pro	Pro	Ala		Pro 3935	Pro
Thr	Thr		Asn 3940	Arg	His	Arg		G1n 3945	Ile	Asp	Arg		Va1 3950	Thr	His
Leu		Ile 3955	Ser	Gly	Leu		Met 3960	Pro	Arg	Gly		A1a 3965	Ile	Asp	Trp
	A1a 3970	Gly	Asn	Val		Trp 3975	Thr	Asp	Ser	Gly	Arg 3980	Asp	Val	Ile	Glu
Va 1 985	Ala	Gln	Met		G1 <i>y</i> 3990	Glu	Asn	Arg		Thr 3995		Ile	Ser		Met 4000
Пe				•					•	ンフフン				•	TUUU

	-	4	1020	-			4	1025				4	1030	Ala	
Asp	_	Thr 1035		Arg			Leu 1040		Gln	-		Ile 1045	Gln	Trp	Pro
	Gly 1050	Leu	Ala	Val		-	His		Glu		Leu 1060	Tyr	Trp	Ala	Asp
A1a 065	Lys				Ile 1070	Gly	Ser	Ile	4	1075					1080
Ile	Val	Ala		Asp 1085	Ser	Lys	Arg		Leu 1090		His	Pro		Ser 1095	Ile
Asp	Val			Asp					Val				Asn 1110	Asn	Arg
Va1		Lys 1115		His								Tyr 1125	Asn	Leu	Thr
		Leu	Ser		Ala	Ser		Val	Val	Leu	Tyr 4140	His	G1n	His	Lys
				Thr								Cys	Glu	Trp	Leu
145				4	1150				4	4155					1160
Cys	Leu	Leu		Pro 4165			Pro		Cys 4170					Gly 4175	Lys
Arg	Leu	•	Asn 4180	-		Cys	Val		Va1		Ser	Pro	Thr 4190	Pro	Pro
Pro	Asp	Ala	Pro	Arg	Pro			Cys	Thr	Leu			Phe	Asn	Gly
		1195					4200					4205			
	Ser 4210	Cys	Phe	Leu		Ala 4215	Arg				Lys 4220	Cys	Arg	Cys	Gln
Pro 225	Arg	Tyr	Thr	-	•		Cys			Asp 4235		Cys	Trp	G1u	Tyr 4240
Cys	His	Asn	-	G1y 4245			Ala				Ser			Pro 4255	Thr
Cys	Arg	-					Thr					Thr		Gln	Va1
Cys				Cys	Ser				Thr	Cys				Gln	Gly
			Gln	Cys	_			Pro	Gly				Asp	Arg	Cys
		Arg	Gln	Cys	Ser	Gly	Phe	Cys	Glu	Asn	Phe	Gly	Thr	Cys	G1n
305	·				4310					4315				-	4320
	Ala	Ala	-	Gly 4325	Ser	Arg	G1n					Val		Phe 4335	
Gly	Pro			Glu	Val	Asn			Ser	Arg	Cys			Gly	Ala

A1a 545

								51/6	35						
Cys		Va1 1355	Asn	Lys	G1n		-	Asp			-	Asn 1365	Cys	Thr	Asp
-			Ala	Pro			Leu	Thr		Пe			Cys	Ser	Asn
G1 y 385	Gly	Ser	Cys					Lys				Glu	-		Cys 1400
Pro	Pro	His			Gly		-	Cys		Glu	Gln	Val	Val	Ser 1415	G1n
Gln	G1n		Gly	His	Met	Ala	Ser	Ile 1425	Leu	Ile	Pro	Leu			Leu
Leu						Ala	Gly	Val	Val	Phe	Trp		_	Arg	Arg
			Ala	•	-	Phe	Gln	His	Gln		Met	Thr	Asn	Gly	Ala
		Va1	Glu	Ile	Gly	Asn	Pro	Thr	Tyr	-		•		•	G1y 1480
Glu	Pro	Asp	•	Val	Gly	Gly	Leu	Leu	Asp	Ala	Asp	Phe	Ala		Asp
Pro	Asp	-						Asn 4505			-		Thr 4510	Leu	Tyr
Met	_	Gly 1515	His	Gly	Ser	_		Ser				Thr 4525	Asp	Glu	Lys
_	G1u 1530	Leu	Leu	Gly	_		Pro	Glu	Asp	Glu		Gly	Asp	Pro	Leu

FIG.6B-14

								52/	65						
GCTAC	CAATO	C AT	CTGG	TCTC	: стс	CAGO	TCC	TTCT	ттст	GC A	AC A	TG G Met 1			55
												GTC Val			103
												GTT Val			151
												GTC Val			199
												GAG G1u 65			247
												AAT Asn			295
												AAT Asn			343
					Gln		Lys	Gly	Pro	Thr		GAA G1u		Lys	391
												GTC Val			439
												AAA Lys 145			487
							His					Leu		CTA Leu	535

FIG.7A-1

53/65 GTA TAC ATT CAG GAT CCC AAA GGA AAT CGC ATC GCA CAA TGG CAG AGT Val Tyr Ile Gln Asp Pro Lys Gly Asn Arg Ile Ala Gln Trp Gln Ser TTC CAG TTA GAG GGT GGC CTC AAG CAA TTT TCT TTT CCC CTC TCA TCA Phe Gln Leu Glu Gly Gly Leu Lys Gln Phe Ser Phe Pro Leu Ser Ser GAG CCC TTC CAG GGC TCC TAC AAG GTG GTG GTA CAG AAG AAA TCA GGT Glu Pro Phe Gln Gly Ser Tyr Lys Val Val Val Gln Lys Lys Ser Gly GGA AGG ACA GAG CAC CCT TTC ACC GTG GAG GAA TTT GTT CTT CCC AAG Gly Arg Thr Glu His Pro Phe Thr Val Glu Glu Phe Val Leu Pro Lys TTT GAA GTA CAA GTA ACA GTG CCA AAG ATA ATC ACC ATC TTG GAA GAA Phe Glu Val Gln Val Thr Val Pro Lys Ile Ile Thr Ile Leu Glu Glu GAG ATG AAT GTA TCA GTG TGT GGC CTA TAC ACA TAT GGG AAG CCT GTC Glu Met Asn Val Ser Val Cys Gly Leu Tyr Thr Tyr Gly Lys Pro Val CCT GGA CAT GTG ACT GTG AGC ATT TGC AGA AAG TAT AGT GAC GCT TCC Pro Gly His Val Thr Val Ser Ile Cys Arg Lys Tyr Ser Asp Ala Ser GAC TGC CAC GGT GAA GAT TCA CAG GCT TTC TGT GAG AAA TTC AGT GGA Asp Cys His Gly Glu Asp Ser Gln Ala Phe Cys Glu Lys Phe Ser Gly CAG CTA AAC AGC CAT GGC TGC TTC TAT CAG CAA GTA AAA ACC AAG GTC Gln Leu Asn Ser His Gly Cys Phe Tyr Gln Gln Val Lys Thr Lys Val TTC CAG CTG AAG AGG AAG GAG TAT GAA ATG AAA CTT CAC ACT GAG GCC Phe Gln Leu Lys Arg Lys Glu Tyr Glu Met Lys Leu His Thr Glu Ala CAG ATC CAA GAA GAA GGA ACA GTG GTG GAA TTG ACT GGA AGG CAG TCC Gln Ile Gln Glu Glu Gly Thr Val Val Glu Leu Thr Gly Arg Gln Ser

FIG.7A-2

54/65 AGT GAA ATC ACA AGA ACC ATA ACC AAA CTC TCA TTT GTG AAA GTG GAC Ser Glu Ile Thr Arg Thr Ile Thr Lys Leu Ser Phe Val Lys Val Asp TCA CAC TTT CGA CAG GGA ATT CCC TTC TTT GGG CAG GTG CGC CTA GTA Ser His Phe Arg Gln Gly Ile Pro Phe Phe Gly Gln Val Arg Leu Val GAT GGG AAA GGC GTC CCT ATA CCA AAT AAA GTC ATA TTC ATC AGA GGA Asp Gly Lys Gly Val Pro Ile Pro Asn Lys Val Ile Phe Ile Arg Gly AAT GAA GCA AAC TAT TAC TCC AAT GCT ACC ACG GAT GAG CAT GGC CTT Asn Glu Ala Asn Tyr Tyr Ser Asn Ala Thr Thr Asp Glu His Gly Leu GTA CAG TTC TCT ATC AAC ACC ACC AAC GTT ATG GGT ACC TCT CTT ACT Val Gln Phe Ser Ile Asn Thr Thr Asn Val Met Gly Thr Ser Leu Thr GTT AGG GTC AAT TAC AAG GAT CGT AGT CCC TGT TAC GGC TAC CAG TGG Val Arg Val Asn Tyr Lys Asp Arg Ser Pro Cys Tyr Gly Tyr Gln Trp GTG TCA GAA GAA CAC GAA GAG GCA CAT CAC ACT GCT TAT CTT GTG TTC Val Ser Glu Glu His Glu Glu Ala His His Thr Ala Tyr Leu Val Phe TCC CCA AGC AAG AGC TIT GTC CAC CTT GAG CCC ATG TCT CAT GAA CTA Ser Pro Ser Lys Ser Phe Val His Leu Glu Pro Met Ser His Glu Leu CCC TGT GGC CAT ACT CAG ACA GTC CAG GCA CAT TAT ATT CTG AAT GGA Pro Cys Gly His Thr Gln Thr Val Gln Ala His Tyr Ile Leu Asn Gly GGC ACC CTG CTG GGG CTG AAG AAG CTC TCC TTT TAT TAT CTG ATA ATG Gly Thr Leu Leu Gly Leu Lys Lys Leu Ser Phe Tyr Tyr Leu Ile Met GCA AAG GGA GGC ATT GTC CGA ACT GGG ACT CAT GGA CTG CTT GTG AAG Ala Lys Gly Gly Ile Val Arg Thr Gly Thr His Gly Leu Leu Val Lys

FIG.7A-3

55/65 CAG GAA GAC ATG AAG GGC CAT TTT TCC ATC TCA ATC CCT GTG AAG TCA Gln Glu Asp Met Lys Gly His Phe Ser Ile Ser Ile Pro Val Lys Ser GAC ATT GCT CCT GTC GCT CGG TTG CTC ATC TAT GCT GTT TTA CCT ACC Asp Ile Ala Pro Val Ala Arg Leu Leu Ile Tyr Ala Val Leu Pro Thr GGG GAC GTG ATT GGG GAT TCT GCA AAA TAT GAT GTT GAA AAT TGT CTG Gly Asp Val Ile Gly Asp Ser Ala Lys Tyr Asp Val Glu Asn Cys Leu GCC AAC AAG GTG GAT TTG AGC TTC AGC CCA TCA CAA AGT CTC CCA GCC Ala Asn Lys Val Asp Leu Ser Phe Ser Pro Ser Gln Ser Leu Pro Ala TCA CAC GCC CAC CTG CGA GTC ACA GCG GCT CCT CAG TCC GTC TGC GCC Ser His Ala His Leu Arg Val Thr Ala Ala Pro Gln Ser Val Cys Ala CTC CGT GCT GTG GAC CAA AGC GTG CTG CTC ATG AAG CCT GAT GCT GAG Leu Arg Ala Val Asp Gln Ser Val Leu Leu Met Lys Pro Asp Ala Glu CTC TCG GCG TCC TCG GTT TAC AAC CTG CTA CCA GAA AAG GAC CTC ACT Leu Ser Ala Ser Ser Val Tyr Asn Leu Leu Pro Glu Lys Asp Leu Thr GGC TTC CCT GGG CCT TTG AAT GAC CAG GAC GAT GAA GAC TGC ATC AAT Gly Phe Pro Gly Pro Leu Asn Asp Gln Asp Asp Glu Asp Cys Ile Asn CGT CAT AAT GTC TAT ATT AAT GGA ATC ACA TAT ACT CCA GTA TCA AGT Arg His Asn Val Tyr Ile Asn Gly Ile Thr Tyr Thr Pro Val Ser Ser ACA AAT GAA AAG GAT ATG TAC AGC TTC CTA GAG GAC ATG GGC TTA AAG Thr Asn Glu Lys Asp Met Tyr Ser Phe Leu Glu Asp Met Gly Leu Lys GCA TTC ACC AAC TCA AAG ATT CGT AAA CCC AAA ATG TGT CCA CAG CTT Ala Phe Thr Asn Ser Lys Ile Arg Lys Pro Lys Met Cys Pro Gln Leu

FIG.7A-4

56/65 CAA CAG TAT GAA ATG CAT GGA CCT GAA GGT CTA CGT GTA GGT TTT TAT Gln Gln Tyr Glu Met His Gly Pro Glu Gly Leu Arg Val Gly Phe Tyr GAG TCA GAT GTA ATG GGA AGA GGC CAT GCA CGC CTG GTG CAT GTT GAA Glu Ser Asp Val Met Gly Arg Gly His Ala Arg Leu Val His Val Glu GAG CCT CAC ACG GAG ACC GTA CGA AAG TAC TTC CCT GAG ACA TGG ATC Glu Pro His Thr Glu Thr Val Arg Lys Tyr Phe Pro Glu Thr Trp Ile TGG GAT TTG GTG GTG GTA AAC TCA GCA GGG GTG GCT GAG GTA GGA GTA Trp Asp Leu Val Val Val Asn Ser Ala Gly Val Ala Glu Val Gly Val ACA GTC CCT GAC ACC ATC ACC GAG TGG AAG GCA GGG GCC TTC TGC CTG Thr Val Pro Asp Thr Ile Thr Glu Trp Lys Ala Gly Ala Phe Cys Leu TCT GAA GAT GCT GGA CTT GGT ATC TCT TCC ACT GCC TCT CTC CGA GCC Ser Glu Asp Ala Gly Leu Gly Ile Ser Ser Thr Ala Ser Leu Arg Ala TTC CAG CCC TTC TTT GTG GAG CTT ACA ATG CCT TAC TCT GTG ATT CGT Phe Gln Pro Phe Phe Val Glu Leu Thr Met Pro Tyr Ser Val Ile Arg GGA GAG GCC TTC ACA CTC AAG GCC ACG GTC CTA AAC TAC CTT CCC AAA Gly Glu Ala Phe Thr Leu Lys Ala Thr Val Leu Asn Tyr Leu Pro Lys TGC ATC CGG GTC AGT GTG CAG CTG GAA GCC TCT CCC GCC TTC CTT GCT Cys Ile Arg Val Ser Val Gln Leu Glu Ala Ser Pro Ala Phe Leu Ala GTC CCA GTG GAG AAG GAA CAA GCG CCT CAC TGC ATC TGT GCA AAC GGG Val Pro Val Glu Lys Glu Gln Ala Pro His Cys Ile Cys Ala Asn Gly CGG CAA ACT GTG TCC TGG GCA GTA ACC CCA AAG TCA TTA GGA AAT GTG Arg Gln Thr Val Ser Trp Ala Val Thr Pro Lys Ser Leu Gly Asn Val

FIG.7A-5

57/65 AAT TTC ACT GTG AGC GCA GAG GCA CTA GAG TCT CAA GAG CTG TGT GGG Asn Phe Thr Val Ser Ala Glu Ala Leu Glu Ser Gln Glu Leu Cys Gly

ACT GAG GTG CCT TCA GTT CCT GAA CAC GGA AGG AAA GAC ACA GTC ATC Thr Glu Val Pro Ser Val Pro Glu His Gly Arg Lys Asp Thr Val Ile

AAG CCT CTG TTG GTT GAA CCT GAA GGA CTA GAG AAG GAA ACA ACA TTC Lys Pro Leu Leu Val Glu Pro Glu Gly Leu Glu Lys Glu Thr Thr Phe

AAC TCC CTA CTT TGT CCA TCA GGT GGT GAG GTT TCT GAA GAA TTA TCC Asn Ser Leu Leu Cys Pro Ser Gly Gly Glu Val Ser Glu Glu Leu Ser

CTG AAA CTG CCA CCA AAT GTG GTA GAA GAA TCT GCC CGA GCT TCT GTC Leu Lys Leu Pro Pro Asn Val Val Glu Glu Ser Ala Arg Ala Ser Val

TCA GTT TTG GGA GAC ATA TTA GGC TCT GCC ATG CAA AAC ACA CAA AAT Ser Val Leu Gly Asp Ile Leu Gly Ser Ala Met Gln Asn Thr Gln Asn

CTT CTC CAG ATG CCC TAT GGC TGT GGA GAG CAG AAT ATG GTC CTC TTT Leu Leu Gln Met Pro Tyr Gly Cys Gly Glu Gln Asn Met Val Leu Phe

GCT CCT AAC ATC TAT GTA CTG GAT TAT CTA AAT GAA ACA CAG CAG CTT Ala Pro Asn Ile Tyr Val Leu Asp Tyr Leu Asn Glu Thr Gln Gln Leu

ACT CCA GAG GTC AAG TCC AAG GCC ATT GGC TAT CTC AAC ACT GGT TAC Thr Pro Glu Val Lys Ser Lys Ala Ile Gly Tyr Leu Asn Thr Gly Tyr

CAG AGA CAG TTG AAC TAC AAA CAC TAT GAT GGC TCC TAC AGC ACC TTT Gln Arg Gln Leu Asn Tyr Lys His Tyr Asp Gly Ser Tyr Ser Thr Phe

GGG GAG CGA TAT GGC AGG AAC CAG GGC AAC ACC TGG CTC ACA GCC TTT Gly Glu Arg Tyr Gly Arg Asn Gln Gly Asn Thr Trp Leu Thr Ala Phe

FIG. 7A-6

				Phe					Ala				ATC Ile	Asp	3223
			Thr					Trp					CAG G1n		3271
		Cys					Gly					Asn	GCC A1a 1090		3319
	Gly					۷a٦					Tyr		ACC Thr		3367
Leu					Leu					Pro			CGC Arg		3415
				Glu					Thr				GGG Gly	Asp	3463
			Val					Leu					TTT Phe		3511
		Asn					Lys					Ser	CTT Leu 1170		3559
	Ala					Asn					Glu		CCT Pro		3607
Pro					Gly					Pro			CCC Pro		3655
	Val			Thr		Tyr			Leu		Tyr		ACG Thr	Ala	 3703

59/65 CCA GCC CCA ACC TCG GAG GAC CTG ACC TCT GCA ACC AAC ATC GTG AAG Pro Ala Pro Thr Ser Glu Asp Leu Thr Ser Ala Thr Asn Ile Val Lys TGG ATC ACG AAG CAG CAG AAT GCC CAG GGC GGT TTC TCC TCC ACC CAG Trp Ile Thr Lys Gln Gln Asn Ala Gln Gly Gly Phe Ser Ser Thr Gln GAC ACA GTG GTG GCT CTC CAT GCT CTG TCC AAA TAT GGA GCC GCC ACA Asp Thr Val Val Ala Leu His Ala Leu Ser Lys Tyr Gly Ala Ala Thr TTT ACC AGG ACT GGG AAG GCT GCA CAG GTG ACT ATC CAG TCT TCA GGG Phe Thr Arg Thr Gly Lys Ala Ala Gln Val Thr Ile Gln Ser Ser Gly ACA TTT TCC AGC AAA TTC CAA GTG GAC AAC AAC AAT CGC CTG TTA CTG Thr Phe Ser Ser Lys Phe Gln Val Asp Asn Asn Asn Arg Leu Leu Leu CAG CAG GTC TCA TTG CCA GAG CTG CCT GGG GAA TAC AGC ATG AAA GTG Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr Ser Met Lys Val ACA GGA GAA GGA TGT GTC TAC CTC CAG ACC TCC TTG AAA TAC AAT ATT Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu Lys Tyr Asn Ile CTC CCA GAA AAG GAA GAG TTC CCC TTT GCT TTA GGA GTG CAG ACT CTG Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly Val Gln Thr Leu CCT CAA ACT TGT GAT GAA CCC AAA GCC CAC ACC AGC TTC CAA ATC TCC Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile Ser CTA AGT GTC AGT TAC ACA GGG AGC CGC TCT GCC TCC AAC ATG GCG ATC Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile GTT GAT GTG AAG ATG GTC TCT GGC TTC ATT CCC CTG AAG CCA ACA GTG Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu Lys Pro Thr Val

FIG.7A-8

60/65	
AAA ATG CTT GAA AGA TCT AAC CAT GTG AGC CGG ACA GAA GTC AGC AGC 427	79
Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu Val Ser Ser	
1400 1405 1410	
AAC CAT GTC TTG ATT TAC CTT GAT AAG GTG TCA AAT CAG ACA CTG AGC 432	27
Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln Thr Leu Ser	L 1
1415 1420 1425	
1713 1720 1723	
TTG TTC TTC ACG GTT CTG CAA GAT GTC CCA GTA AGA GAT CTC AAA CCA 43	75
Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val Arg Asp Leu Lys Pro	
1430 1435 1440	
GCC ATA GTG AAA GTC TAT GAT TAC TAC GAG ACG GAT GAG TTT GCA ATC 44:	23
Ala Ile Val Lys Val Tyr Asp Tyr Tyr Glu Thr Asp Glu Phe Ala Ile	
1445 1450 1455 1460	
COT OAG TAG AAT COT CCT TCC ACC AAA CAT CTT COA AAT CCT TCAACACOA	~ .
GCT GAG TAC AAT GCT CCT TGC AGC AAA GAT CTT GGA AAT GCT TGAAGACCA 44	/4
Ala Glu Tyr Asn Ala Pro Cys Ser Lys Asp Leu Gly Asn Ala	
1465 1470 1	
CAAGGCTGAA AAGTGCTTTG CTGGAGTCCT GTTCTCTGAG CTCCACAGAA GACACGTGTT 45.	34
TTTGTATCTT TAAAGACTTG ATGAATAAAC ACTTTTTCTG GTC 45	

FIG.7A-9

Ser Val Ser Gly Lys Pro Gln Tyr Met Val Leu Val Pro Ser Leu Leu His Thr Glu Thr Thr Glu Lys Gly Cys Val Leu Leu Ser Tyr Leu Asn Glu Thr Val Thr Val Ser Ala Ser Leu Glu Ser Val Arg Gly Asn Arg Ser Leu Phe Thr Asp Leu Glu Ala Glu Asn Asp Val Leu His Cys Val Ala Phe Ala Val Pro Lys Ser Ser Ser Asn Glu Glu Val Met Phe Leu Thr Val Gln Val Lys Gly Pro Thr Gln Glu Phe Lys Lys Arg Thr Thr Val Met Val Lys Asn Glu Asp Ser Leu Val Phe Val Gln Thr Asp Lys Ser Ile Tyr Lys Pro Gly Gln Thr Val Lys Phe Arg Val Val Ser Met Asp Glu Asn Phe His Pro Leu Asn Glu Leu Ile Pro Leu Val Tyr Ile Gln Asp Pro Lys Gly Asn Arg Ile Ala Gln Trp Gln Ser Phe Gln Leu Glu Gly Gly Leu Lys Gln Phe Ser Phe Pro Leu Ser Ser Glu Pro Phe Gln Gly Ser Tyr Lys Val Val Val Gln Lys Lys Ser Gly Gly Arg Thr Glu His Pro Phe Thr Val Glu Glu Phe Val Leu Pro Lys Phe Glu Val Gln Val Thr Val Pro Lys Ile Ile Thr Ile Leu Glu Glu Glu Met Asn Val Ser Val Cys Gly Leu Tyr Thr Tyr Gly Lys Pro Val Pro Gly His Val Thr Val Ser Ile Cys Arg Lys Tyr Ser Asp Ala Ser Asp Cys His Gly Glu Asp Ser Gln Ala Phe Cys Glu Lys Phe Ser Gly Gln Leu Asn Ser His Gly Cys Phe Tyr Gln Gln Val Lys Thr Lys Val Phe Gln Leu Lys Arg Lys Glu Tyr Glu Met Lys Leu His Thr Glu Ala Gln Ile Gln Glu Glu Gly Thr Val Val Glu Leu Thr Gly Arg Gln Ser Ser Glu Ile

FIG. 7B-1

Thr Arg Thr Ile Thr Lys Leu Ser Phe Val Lys Val Asp Ser His Phe Arg Gln Gly Ile Pro Phe Phe Gly Gln Val Arg Leu Val Asp Gly Lys Gly Val Pro Ile Pro Asn Lys Val Ile Phe Ile Arg Gly Asn Glu Ala Asn Tyr Tyr Ser Asn Ala Thr Thr Asp Glu His Gly Leu Val Gln Phe Ser Ile Asn Thr Thr Asn Val Met Gly Thr Ser Leu Thr Val Arg Val Asn Tyr Lys Asp Arg Ser Pro Cys Tyr Gly Tyr Gln Trp Val Ser Glu Glu His Glu Glu Ala His His Thr Ala Tyr Leu Val Phe Ser Pro Ser Lys Ser Phe Val His Leu Glu Pro Met Ser His Glu Leu Pro Cys Gly His Thr Gln Thr Val Gln Ala His Tyr Ile Leu Asn Gly Gly Thr Leu Leu Gly Leu Lys Lys Leu Ser Phe Tyr Tyr Leu Ile Met Ala Lys Gly Gly Ile Val Arg Thr Gly Thr His Gly Leu Leu Val Lys Gln Glu Asp Met Lys Gly His Phe Ser Ile Ser Ile Pro Val Lys Ser Asp Ile Ala Pro Val Ala Arg Leu Leu Ile Tyr Ala Val Leu Pro Thr Gly Asp Val Ile Gly Asp Ser Ala Lys Tyr Asp Val Glu Asn Cys Leu Ala Asn Lys Val Asp Leu Ser Phe Ser Pro Ser Gln Ser Leu Pro Ala Ser His Ala His Leu Arg Val Thr Ala Ala Pro Gln Ser Val Cys Ala Leu Arg Ala Val Asp Gln Ser Val Leu Leu Met Lys Pro Asp Ala Glu Leu Ser Ala Ser Ser Val Tyr Asn Leu Leu Pro Glu Lys Asp Leu Thr Gly Phe Pro Gly Pro Leu Asn Asp Gln Asp Asp Glu Asp Cys Ile Asn Arg His Asn Val Tyr Ile Asn Gly Ile Thr Tyr Thr Pro Val Ser Ser Thr Asn Glu Lys Asp Met Tyr Ser Phe Leu Glu Asp Met Gly Leu Lys Ala Phe Thr

Asn Ser Lys Ile Arg Lys Pro Lys Met Cys Pro Gln Leu Gln Gln Tyr Glu Met His Gly Pro Glu Gly Leu Arg Val Gly Phe Tyr Glu Ser Asp Val Met Gly Arg Gly His Ala Arg Leu Val His Val Glu Glu Pro His Thr Glu Thr Val Arg Lys Tyr Phe Pro Glu Thr Trp Ile Trp Asp Leu Val Val Val Asn Ser Ala Gly Val Ala Glu Val Gly Val Thr Val Pro Asp Thr Ile Thr Glu Trp Lys Ala Gly Ala Phe Cys Leu Ser Glu Asp Ala Gly Leu Gly Ile Ser Ser Thr Ala Ser Leu Arg Ala Phe Gln Pro Phe Phe Val Glu Leu Thr Met Pro Tyr Ser Val Ile Arg Gly Glu Ala Phe Thr Leu Lys Ala Thr Val Leu Asn Tyr Leu Pro Lys Cys Ile Arg Val Ser Val Gln Leu Glu Ala Ser Pro Ala Phe Leu Ala Val Pro Val Glu Lys Glu Gln Ala Pro His Cys Ile Cys Ala Asn Gly Arg Gln Thr Val Ser Trp Ala Val Thr Pro Lys Ser Leu Gly Asn Val Asn Phe Thr Val Ser Ala Glu Ala Leu Glu Ser Gln Glu Leu Cys Gly Thr Glu Val Pro Ser Val Pro Glu His Gly Arg Lys Asp Thr Val Ile Lys Pro Leu Leu Val Glu Pro Glu Gly Leu Glu Lys Glu Thr Thr Phe Asn Ser Leu Leu Cys Pro Ser Gly Gly Glu Val Ser Glu Glu Leu Ser Leu Lys Leu Pro Pro Asn Val Val Glu Glu Ser Ala Arg Ala Ser Val Ser Val Leu Gly Asp Ile Leu Gly Ser Ala Met Gln Asn Thr Gln Asn Leu Leu Gln Met Pro Tyr Gly Cys Gly Glu Gln Asn Met Val Leu Phe Ala Pro Asn Ile Tyr Val Leu Asp Tyr Leu Asn Glu Thr Gln Gln Leu Thr Pro Glu Val Lys Ser Lys Ala Ile Gly Tyr Leu Asn Thr Gly Tyr Gln Arg Gln

Lys Val Thr	Gln Ala Gly 1250	Gln Leu 1235 Lys	Asn 1220 His Ala	1205 Ala Ala Ala	Gln Leu Gln	Gly Ser Val	Gly Lys 1240 Thr	Thr Phe 1225 Tyr Ile	Asn 1210 Ser Gly Gln	Ile Ser Ala Ser	Thr Ala Ser 1260	GIn Thr 1245 Gly	Asp 1230 Phe Thr	Ile 1215 Thr	Val Arg Ser
Lys Val Thr	Gln Ala Gly	Gln Leu 1235	Asn 1220 His	1205 Ala Ala	Gln Leu Gln	Gly Ser Val	Ala Gly Lys 1240	Thr Phe 1225 Tyr	Asn 1210 Ser Gly	Ile Ser Ala Ser	Thr Ala Ser	G1n Thr 1245	Asp 1230 Phe	Ile 1215 Thr Thr	Val Arg
Lys	Gln Ala	Gln Leu	Asn 1220	1205 Ala	Gln	Gly Ser	Ala Gly Lys	Thr : Phe 1225	Asn 1210 Ser	Ile Ser	Thr Ala	G1n Thr	Asp 1230	Ile 1215 Thr	Val
		Gln	Asn	1205 Ala	GIn	Gly	Ala Gly	Thr : Phe	Asn 1210 Ser	Ile Ser	Thr	GIn	Asp	Ile 1215 Thr	
			•	1205			Ala	Thr	Asn 1210	Ile				Ile 1215	
T1	Ser	13 111	A	1	Tl	C					1/-7	1	т		T1
185			1,51		1190.		, u		LCG.	 1195	, (, a	u			1200
	1170 Thr		Tvr			1175 1 eu				Thr		GIn	Pro	Ala	Pro
	Val	Gly		Phe	Tyr	Glu	Pro	Gln	Ala		Ser		G1u	Val	Glu
Lys		Asp	Asn	Ser	Val	His	Trp	Glu	Arg	Pro	Gln			Lys	Ala
Gln	Asp	_				Val					Asn		Glu 1150	Ala	Val
vaı	ıyr	ınr		A1a L125								Leu		Gly 1135	ASN
105				1	1110				1	1115				1	120
			Δla			1095 Thr			Glu		1100 Asp	His	G1 v	Ser	His
Ile			Thr							Arg		L085 A1a	Leu	Phe	Cys
G1u	-	Glu	Val	Thr	Leu	Ser	Ala	Tyr				Ala		Leu	Glu
Phe	Arg			Gly	Ser		Leu					Lys	G1y L070	Gly	Val
	Gln	Ala		Ile	Trp		Ser	Gln	Arg	Gln		Asp	Asn	G1 y L055	Cys
	Phe	Ala	Gln	Ala	Arg	Ala	Tyr	Пe	Phe	Пe	Asp		Ala	His 1	11e .040
				G1n	Gly	Asn	Thr	Trp		Thr		Phe	Va1	Leu	Lys
T		1yr 995				Asp 1						.005	Gly	Glu	Arg

65/65 Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile Ser Leu Ser Val 1340 1335 1330 Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile Val Asp Val 345 1350 1355 1360 Lys Met Val Ser Gly Phe Ile Pro Leu Lys Pro Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln Thr Leu Ser Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val Arg Asp Leu Lys Pro Ala Ile Val 1415 1420 1410 Lys Val Tyr Asp Tyr Tyr Glu Thr Asp Glu Phe Ala Ile Ala Glu Tyr 425 1430 1435 1440

FIG.7B-5

1450

Asn Ala Pro Cys Ser Lys Asp Leu Gly Asn Ala

1445